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**by**

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# Classification of Quadrature Amplitude Modulated (QAM) Signals via Sequential Probability Ratio Test (SPRT)\*

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## Abstract

A novel approach to automatic classification of quadrature amplitude modulated (QAM) signals is presented in this research. Modulation classification has been traditionally treated as a hypothesis test problem with input signals of a fixed sample size. By formulating it as a variable sample size test problem, we propose a new classification algorithm based on the sequential probability ratio test (SPRT). It is demonstrated that the new approach has several important merits, including ease of error rate control, lower computational complexity and lower decision delay.

## 1 Introduction

Automatic modulation classification is an important research problem in the receiver design for non-cooperative communication systems. The modulation classifier usually serves as a preprocessing unit for monitoring or interception systems. When the modulation scheme of a received signal is recognized, an appropriate demodulator can be selected to recover the information. The automatic modulation classification technique can also be applied to the control of radio frequency bands. Such a control task has so far been done by human operators. However, it becomes more difficult due to higher signal densities over a fixed bandwidth in recent years.

The difficulty of this problem is due to the uncertainty of received signals. First, the transmitted information symbol sequence is unknown. Usually, it is reasonable to assume that the statistics of source encoder outputs comply with the maximum entropy principle so that the representation of information is efficient. The second difficulty comes from the channel distortion such as noise, interferences, etc. The additive white Gaussian noise (AWGN) is the most commonly used channel

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model because of its simple analytical results and well known properties in the long history of detection/estimation theory. The third difficulty is that some communication parameters such as the reference phase might not be known in advance. These parameters could be essential before we can further acquire the modulation scheme. Finally, the classification problem can be even more difficult when only a short period of data record is available. This happens in cases such as signal interception and RF band control where the received signal is present only over a very short time period.

The problem of modulation classification consists of two steps: feature extraction from the received waveform and classification according to the extracted features. Several features have been used to characterize different modulation schemes, including the power spectral density [14], [18], bispectrum [14], time-varying frequency [5], [11], [12], [18], [19], [21], time-varying phase [5], [11], [21], [23], [29], [33], time-varying envelope [1], [9], [11], [18] and regenerated spectral lines [24], [27].

Two approaches to obtain the statistics of extracted features have been reported in previous work. The first approach derives the exact statistical descriptions of received signals under certain given assumptions [1], [8], [9], [20], [24], [29], [33]. This approach is often motivated by the receiver design of cooperative communication systems. It gives more insights into the cause-effect relationship among the performance and various uncertainty factors. An exact analysis of feature statistics is possible by imposing the synchronization assumptions, and the optimality can be theoretically guaranteed. The challenging part of this approach is the derivation of test statistics based on a set of reasonable assumptions. The second approach obtains the test statistics empirically rather than analytically, i.e. to design the classifier by using training data. Classifiers using features such as the periodogram, the bispectrum and the histograms of local frequency or phase estimates, have been reported in [5], [11], [12], [14], [23], [27]. This approach is more realistic in the sense that it requires less a priori knowledge of received signals. However, the corresponding theoretical performance bound is very difficult to analyze.

Modulation classification has been traditionally formulated as a hypothesis test problem with a fixed size of samples, i.e. making decisions based on a fixed amount of received data. Classifiers using the fixed-sample-size test (FSST), which often operate in a batch mode, are suitable for bursty or packet type of data, For other applications, such as the transmission of continuum type data, it is more natural to formulate the problem as a sequential process since received data are collected in a

temporal order and their amount can be quite large. In this context, a more appropriate optimality is the least amount of data required for decision making while the classification performance such as the error probability meets a certain requirement.

This work focuses on the classification of signals with the quadrature amplitude modulation (QAM), which forms an important family of digital modulation schemes. We formulate this modulation classification problem as a variable sample size test problem, and propose a new classification algorithm based on the sequential probability ratio test (SPRT). It will be demonstrated that the new approach has several important merits, including ease of error rate control, lower computational complexity and lower decision delay. The derivation of a reference-phase-free test for QAM classification has been an important problem [20], [24]. Three different techniques, i.e. the transform, the generalized likelihood and the averaging likelihood methods, are presented in the paper to resolve the problem of reference phase uncertainty.

This paper is organized as follows. In Sections 2 and 3, we examine modulation classification algorithms using FSST and SPRT, respectively. In Section 4, three techniques are discussed to derive the reference phase invariant classifiers. Experimental results are shown in Section 5 and concluding remarks are presented in Section 6.

## 2 Fixed-Sample-Size Test (FSST)

In this section, we first formulate the problem, and then review the well known fixed-sample-size test.

### 2.1 Problem Formulation

We assume that the received waveform  $r(t)$  is a QAM modulated signal  $x(t; \vec{s}_N, k, \vec{p})$  buried in AWGN  $n(t)$  with zero mean and two-sided power spectral density of  $N_0$  W/Hz, where  $\vec{s}_N$  represents the symbol sequence  $(s_0, s_1, \dots, s_{N-1})$ ,  $N$  is the total number of received symbols,  $k$  indicates a different modulation format and  $\vec{p}$  denotes a vector of communication parameters. Therefore, the received  $N$  symbol-periods waveform can be written as

$$r(t) = x(t; \vec{s}_N, k, \vec{p}) + n(t), \quad 0 \leq t \leq NT, \quad (1)$$

where  $T$  is the symbol duration. Here we assume that symbol timing is synchronized so that the signal is received at the beginning of each symbol period.

If the transmitted signal waveform  $x(t; \vec{s}_N, k, \vec{p})$  is known, we can write down the a posteriori probability of the transmitted signal waveform based on the received signal  $r(t)$  by using Bayes' rule as

$$P(x(t; \vec{s}_N, k, \vec{p})|r(t)) = \frac{P(x(t; \vec{s}_N, k, \vec{p}))P(r(t)|x(t; \vec{s}_N, k, \vec{p}))}{P(r(t))}.$$

By assuming that  $\vec{p}$  is known or can be measured accurately from preprocessors, we can drop it for discussion in Sections 2 and 3. (Three methods will be described to handle the case of unknown  $\vec{p}$ , especially in resolving reference phase uncertainty, in Section 4.) We further assume that there is no preference for any particular symbol sequence or modulation format so that  $P(x(t; \vec{s}, k, \vec{p}))$  and  $P(r(t))$  are simply normalization terms which are independent of  $\vec{s}$  or  $k$ . Then, the a posteriori probability of  $x(t; \vec{s}_N, k)$  can be expressed as the following well-known form [31]

$$P(x(t; \vec{s}_N, k)|r(t)) = C \exp \left\{ \frac{-1}{2N_0} \int_0^{NT} [r(t) - x(t; \vec{s}_N, k)]^2 dt \right\}. \quad (2)$$

A QAM signal  $x(t; \vec{s}_N, k)$  can be written as: for  $0 \leq t \leq NT$ ,

$$x(t; \vec{s}_N, k) = X_I(t; \vec{s}_N, k) \cos(2\pi f_c t - \theta_c) + X_Q(t; \vec{s}_N, k) \sin(2\pi f_c t - \theta_c), \quad (3)$$

where  $f_c$  and  $\theta_c$  denote the carrier frequency and the carrier reference phase, respectively, and

$$\begin{aligned} X_I(t; \vec{s}_N, k) &= \sum_{i=0}^N x_I(i; k) p(t - iT), \\ X_Q(t; \vec{s}_N, k) &= \sum_{i=0}^N x_Q(i; k) p(t - iT), \end{aligned}$$

are the information bearing waveforms for I- and Q-channels, and where  $p(t)$  is the spectrum shaping pulse function which is assumed to be known and absorbed into the communication parameter vector  $\vec{p}$ . In particular, we let  $p(t)$  be a rectangular unit pulse function

$$p(t) = \begin{cases} 1 & \text{if } 0 \leq t \leq T, \\ 0 & \text{otherwise.} \end{cases}$$

The I-, Q-channel sample sequences  $\{x_I(i; k); i = 1, 2, \dots\}$  and  $\{x_Q(i; k); i = 1, 2, \dots\}$  can be represented by  $\{(x_I(i; k), x_Q(i; k)) \in \mathcal{S}_k; i = 1, 2, \dots\}$ , where  $\mathcal{S}_k \triangleq \{(s_I(m; k), s_Q(m; k)); m = 1, \dots, M\}$  denotes the constellation or the symbol set of the modulation type  $k$ . The resulting two dimensional plot of these  $M$  vectors  $(s_I(m; k), s_Q(m; k))$ ,  $m = 1, \dots, M$  is called the constellation or signal space diagram, where each point in the diagram is called a constellation point. Examples are given in Fig. 1 to illustrate different constellation diagrams for QAM signals.

By imposing the assumptions of independent and identically distributed (i.i.d.) symbol sequences and an equal probable symbol set, we can derive the following equations from (2)

$$P(x(t; k)|r(t)) = C' \prod_{i=1}^N p(I_i, Q_i|k). \quad (4)$$

Note that  $(I_i, Q_i)$  is the  $i$ th sample of the I-, Q-channel outputs and

$$p(I_i, Q_i|k) = \frac{1}{M} \sum_{m=1}^M \frac{1}{2\pi\sigma_T^2} \exp \left\{ \frac{-1}{2\sigma_T^2} [(I_i - s_I(m; k))^2 + (Q_i - s_Q(m; k))^2] \right\}, \quad (5)$$

where  $(s_I(m; k), s_Q(m; k))$  is the  $m$ -th constellation point of the  $k$ -th modulation scheme and  $\sigma_T^2 = \frac{N_0 T}{2}$ .

## 2.2 MAP Classifier

Suppose that there are  $K$  possible modulation schemes in association with an  $N$ -symbol received waveform. Let us construct a multihypothesis test by associating each of the  $K$  hypotheses  $H_1, H_2, \dots, H_K$  with one modulation scheme. The a posteriori probability  $P(x(t; k)|r(t))$  under  $H_k$  is specified by Eq. (4). Based on the maximum a posteriori (MAP) principle, one can minimize the total (or average) error probability

$$P_e = \sum_{i=1}^K P(H_i) \epsilon_i,$$

where the individual decision error  $\epsilon_i$  is defined by

$$\epsilon_i = \sum_{j \neq i} \text{Prob}(\text{decided on } H_j | H_i \text{ is true}),$$

by deciding on  $H_{\hat{k}}$  whenever its a posteriori probability is the largest. In other words, the modulation type  $\hat{k}$  is chosen by using the following criterion

$$\hat{k} = \arg \max_{k=1}^K P(x(t; k)|r(t)).$$

Note that the MAP classifier makes decision based on the probability calculation using a fixed number  $N$  of received symbols.

## 3 Variable-Sample-Size Test with SPRT

A statistical test that uses a random number of samples is called the *sequential* test [15], [16], [25]. One important advantage of the sequential test is the flexibility in controlling the individual error

probability  $\epsilon_i$  for a multihypothesis test problem. The MAP test discussed above minimizes the total error probability, but has no control over the individual error probability conditioned on a given hypothesis. In a binary hypothesis test, the Neyman-Pearson test maximizes the correct probability of one hypothesis (i.e. the detection probability) while keeping the false alarm probability under a certain level. Although it is possible to generalize the Neyman-Pearson test to the multihypothesis case, this technique is not widely used in practice [31]. In this section, we propose the use of the sequential probability ratio test (SPRT) for QAM classification for binary as well as multihypothesis cases. With the sequential test, we can keep performing the test with more observations until a certain performance or stopping criterion is achieved.

### 3.1 SPRT for Binary Hypothesis

A binary sequential test [25] can be stated as follows. By assuming that the observed sequence of random variables  $X_1, X_2, X_3, \dots$  are i.i.d., we would like to determine which distribution ( $P_1$  or  $P_2$ ) the observed sequence comes from. In mathematics, this test can be written as

$$\begin{aligned} H_1 & : X_k \sim P_1, \quad k = 1, 2, \dots, \\ H_2 & : X_k \sim P_2, \quad k = 1, 2, \dots, \end{aligned}$$

where the notation “ $\sim$ ” denotes “obtained from a certain distribution”. A sequential test is characterized by a set of stopping rules and a set of decision rules, denoted by  $\bar{\Phi} = \{\Phi_1, \Phi_2, \dots\}$  and  $\bar{\Sigma} = \{\Sigma_1, \Sigma_2, \dots\}$ , respectively. The stopping rule at time  $n$  tells us whether we should stop the experiment with observations  $X_1, \dots, X_n$  or continue the experiment for one more additional observation  $X_{n+1}$ . The decision rule performs the hypothesis test based on all available data. Note that FSST of size  $n$  is in fact a special case of a sequential test, namely, it stops at  $n$ .

The sequential probability ratio test (SPRT) provides a very effective sequential test and is defined below. With SPRT, we compute the likelihood ratio

$$\lambda(X_1, X_2, \dots, X_n) = \prod_{k=1}^n \frac{P_2(X_k)}{P_1(X_k)}$$

based on samples  $\{X_1, X_2, \dots, X_n\}$  and compare this ratio with threshold values  $a$  and  $b$  ( $0 \leq a < b < \infty$ ). The test continues until the ratio  $\lambda(X_1, X_2, \dots, X_n)$  falls outside  $(a, b)$ , and the decision rule is

$$\begin{aligned} \text{if } \lambda(X_1, X_2, \dots, X_n) < a & \longrightarrow H_1, \\ \text{if } \lambda(X_1, X_2, \dots, X_n) > b & \longrightarrow H_2, \end{aligned}$$

To explain SPRT better, we consider an example of classifying BPSK/QPSK and show four snapshots in Fig. 2 conditioned on that the source signals are actually BPSK modulated. The x- and y-coordinates of the figure represent the time and the log likelihood ratio value, respectively, and the two parallel lines in each snapshot represent the decision boundaries of SPRT. The four sample curves provides different variations of the log likelihood ratio along the time axis. SPRT will not stop until the log likelihood ratio hits one of the two decision boundaries. If the upper (or lower) boundary is hit first, we decide on BPSK (or QPSK). Time required for decision making depends on the data statistics as well as decision boundaries. If input data are clearly in favor of one of the two hypotheses or decision boundaries are close to zero, it requires a smaller number of samples (or, equivalently, shorter time delay) in decision making. On the contrary, if input data are more uncertain or decision boundaries are away from zero, more time is needed to reach a decision.

SPRT with likelihood ratio  $\lambda$  and interval  $(a, b)$  is usually denoted by  $\text{SPRT}(a, b, \lambda)$ . The following Wald-Wolfowitz theorem [32] characterizes the optimality of SPRT.

**Proposition 1 Wald-Wolfowitz Theorem**

Consider  $\text{SPRT}(a, b, \lambda)$  with stopping rule set  $\bar{\Phi}_0$  and decision rule set  $\bar{\Sigma}_0$ . Let  $(\bar{\Phi}, \bar{\Sigma})$  be the corresponding sets of any other sequential test such that

$$\begin{aligned}\epsilon_1(\bar{\Phi}, \bar{\Sigma}) &\leq \epsilon_1(\bar{\Phi}_0, \bar{\Sigma}_0) \\ \epsilon_2(\bar{\Phi}, \bar{\Sigma}) &\leq \epsilon_2(\bar{\Phi}_0, \bar{\Sigma}_0)\end{aligned}$$

where  $\epsilon_1$  (or  $\epsilon_2$ ) is the error decision probability given that  $H_1$  (or  $H_2$ ) is true. Then,

$$\begin{aligned}E\{N(\bar{\Phi})|H_1\} &\geq E\{N(\bar{\Phi}_0)|H_1\} \\ E\{N(\bar{\Phi})|H_2\} &\geq E\{N(\bar{\Phi}_0)|H_2\},\end{aligned}$$

where  $N(\bar{\Phi})$  is the sample size (stopping time) according to the stopping rule  $\bar{\Phi}$ .

This theorem says that on the average SPRT requires the minimum amount of samples to achieve a given level of performance (i.e. the error probability). Thus, the average sample size of SPRT is not greater than that of FSST with the same performance.

Furthermore, we can control the decision interval  $(a, b)$  to adjust the error probabilities of SPRT. Wald [32] showed that as long as the prescribed error probabilities  $\epsilon_1$  and  $\epsilon_2$  are reasonably small, we can choose

$$a = \epsilon_2/(1 - \epsilon_1) \approx \epsilon_2, \quad \text{and} \quad b = (1 - \epsilon_2)/\epsilon_1 \approx 1/\epsilon_1.$$



They are called the Wald boundaries.

### Average Stopping Time

The average stopping time of SPRT can be estimated by using the following corollary [25].

#### Proposition 2 Corollary to Wald's Identity

Consider a binary hypothesis test

$$\begin{aligned} H_1 &: X_k \sim P_1, \quad k = 1, 2, \dots, \\ H_2 &: X_k \sim P_2, \quad k = 1, 2, \dots, \end{aligned}$$

where  $X_1, X_2, X_3, \dots$  are independent and identically distributed (i.i.d.) solved by SPRT( $a, b, \lambda$ ) with

$$\lambda(X_1, X_2, \dots, X_n) = \prod_{k=1}^n \frac{P_2(X_k)}{P_1(X_k)}.$$

Then, the average stopping time for  $H_1$  and  $H_2$  can approximated as

$$\begin{aligned} E\{N|H_1\} &\approx \frac{1}{\mu_1} \left[ (1 - \epsilon_1) \log \frac{\epsilon_2}{1 - \epsilon_1} + \epsilon_1 \log \frac{1 - \epsilon_2}{\epsilon_1} \right], \\ E\{N|H_2\} &\approx \frac{1}{\mu_2} \left[ \epsilon_2 \log \frac{\epsilon_2}{1 - \epsilon_1} + (1 - \epsilon_2) \log \frac{1 - \epsilon_2}{\epsilon_1} \right], \end{aligned}$$

where  $\epsilon_1 = P(H_2|H_1)$ ,  $\epsilon_2 = P(H_1|H_2)$  and

$$\mu_j = E\{\log \lambda(X)|H_j\} \neq 0, \quad \text{for } j = 1, 2.$$

The above approximations can be further simplified to be

$$E\{N|H_1\} \approx \frac{\log \epsilon_2}{\mu_1} \quad \text{and} \quad E\{N|H_2\} \approx -\frac{\log \epsilon_1}{\mu_2} \quad (6)$$

for reasonably small  $\epsilon_1$  and  $\epsilon_2$ .

### Efficiency

Efficiency of SPRT relative to FSST is defined as the ratio of their average sample sizes with respect to the same error level  $\epsilon_j$  [17], i.e.

$$R_j = \frac{n(\epsilon_j)}{E\{N|H_j\}} \quad \text{for } j = 1, 2, \quad (7)$$

where  $n(\epsilon_j)$  is the number of samples required by FSST to achieve the error level  $\epsilon_j$  when the hypothesis  $H_j$  is true. Let us use  $S_n$  to denote the log likelihood ratio of i.i.d. distributed sequence  $X_1, X_2, \dots, X_n$ :

$$S_n = \sum_{i=1}^n \log \frac{P_2(X_i)}{P_1(X_i)} = \sum_{i=1}^n \log \lambda(X_i).$$

The error probability for  $H_j$  is given by  $\epsilon_j = P(S_n \geq 0 | H_j)$ . Assume  $n \gg 1$ , we can apply the central limiting theorem to  $S_n$  so that

$$\epsilon_j \approx Q\left(\frac{-\sqrt{n}\mu_j}{\sigma_j}\right),$$

where

$$Q(x) = \frac{1}{\sqrt{2\pi}} \int_x^\infty e^{-\frac{y^2}{2}} dy, \quad \mu_j = E\{\log \lambda(X) | H_j\}, \quad \sigma_j^2 = E\{(\log \lambda(X))^2 | H_j\} - \mu_j^2.$$

We may further simplify  $Q(x)$  as [28]:

$$Q(x) \approx \frac{1}{\sqrt{2\pi x}} e^{-\frac{x^2}{2}} \quad \text{for } x \gg 0$$

so that the log of error probability can be approximated by

$$\log \epsilon_j \approx -n \frac{\mu_j^2}{2\sigma_j^2} \quad \text{for } \epsilon_j \ll 1.$$

Therefore, the sample size required to achieve the error level of  $\epsilon_j$  via FSST is

$$n(\epsilon_j) \approx -\frac{2\sigma_j^2}{\mu_j^2} \log \epsilon_j. \quad (8)$$

By plugging (8) and (6) into (7), we obtain the efficiency of SPRT as

$$R_j = -\frac{2\sigma_j^2}{\mu_j} \quad \text{for } j = 1, 2.$$

Thus, efficiency depends on statistics  $\log \lambda(X)$  up to the second order.

### Decision Boundaries

The standard SPRT by Wald compares the likelihood ratio with two parallel boundary lines so that one of the hypothesis is accepted whenever the likelihood ratio reaches one of the boundaries. This condition is illustrated in Fig. 3(a). Although Wald's SPRT is optimal in the sense that it requires the smallest amount of samples for decision making with a given individual error probability, there have been criticisms and modifications of Wald's SPRT [10], [30]. Although it is proved that the sample size is finite with probability one, the size can be too long to be practical. In practice, we may be forced to stop even though the stopping criterion has not been met yet, and the resulting performance is not as good as Wald's SPRT. This is known as *truncated SPRT* as depicted in Fig. 3(b), where we show an example of SPRT truncated at  $n_T$  samples and  $C$  denotes the threshold value for decision making after  $n_T$  samples are observed. Another example proposed

in [2] is the converging boundaries as shown in Fig. 3(c), where decision boundaries are dynamically changing with time. This kind of decision strategy makes it easier to reach a conclusion for longer observation time by relaxing the requirement of making correct decisions. Figs. 3 (d) and (e) are methods proposed by Read [26] and Baruah and Bhattacharjee [6], respectively. Read suggested to apply SPRT only after  $n_0$  samples. Decision boundaries in Fig. 3 (e) are in fact a combination of boundaries in Figs. 3 (c) and (d) In our experiments given in Section 5, we follow Read's work as shown in Fig. 3(d).

Tantaratana [30] reviewed several decision boundaries to reduce the average sample size for a non-perfect statistical model [30]. The general approach is to combine the fixed-sample-size and the sequential tests so that the sequential test is invoked after a fixed amount of data have been collected. The purpose of this mixture design is to maintain the performance at least close to FSST when the parameters are mismatched but still keep average sample size as small as Wald's SPRT when parameters are correctly tuned.

Although the concept of truncation is simple and practical, optimalities on decision boundary design, error control, expected sample size, etc. are technically challenging and has been studied for only very limited cases. No general conclusion such as the Wald's theorem has been reported. SPRT requires accurate description of hypothesized test statistics. In practice the signal distribution may not be known exactly and can only be obtained by estimation. This may increase the approximation errors of the error probability and the expected sample size. To take our problem as an example, the distributions of received signals are parameterized by some unknown parameters such as the carrier phase, which might have to be estimated from the data. The estimation error might cause early stopping of the test and, therefore, degrade the predicted performance.

### 3.2 Multihypothesis Test

As mentioned before, one of the major drawback of FSST is that even though it provides the solution to the lowest *average* error probability, there is no assurance in minimizing the *individual* error probability. SPRT was derived for the binary hypothesis case in the previous subsection, and the relationship between individual error probabilities and Wald's boundaries provides important insights into the multiple hypothesis problem. Several multihypothesis sequential tests have been proposed in the past. Most of them divide the test into a couple of binary tests. We may classify them into [16] (1) the positive approach, which considers simultaneously all possible binary hy-

pothesis tests, and (2) the negative approach, which eliminates the unlikely hypothesis along the process until there is only one left. Another way to classify different approaches is to see how the alternative hypothesis is constructed by using binary hypothesis tests. There are three approaches to represent the test statistics of alternative hypothesis: (1) maximum-likelihood, (2) geometric mean and (3) algebraic mean. These approaches are explained below.

### Positive Approach

The Armitage test [3], [16] generalizes the binary to M-ary hypothesis test by constructing pair-wise SPRT for all hypotheses. For example, if there are  $m$  hypotheses  $H_1, H_2, \dots, H_m$  and their corresponding distribution functions are  $P_1, P_2, \dots, P_m$ , respectively. This approach stops the test at  $n$  samples and accepts  $H_i$  if

$$\prod_{k=1}^n \frac{P_i(X_k)}{P_j(X_k)} > b_{i,j}, \quad \text{for all } j \neq i,$$

where  $X_1, X_2, \dots, X_n$  are i.i.d. samples. The decision error probability  $\epsilon_i$  given that hypothesis  $H_i$  is true can be controlled by adjusting  $b_{i,j}$  and it is bounded by

$$\epsilon_i = \sum_{j \neq i} \epsilon_{i,j} \leq \sum_{j \neq i} \frac{1}{b_{i,j}},$$

where  $\epsilon_{i,j}$  is the error probability of claiming  $H_j$  given that hypothesis  $H_i$  is true.

### Negative Approach

The negative approach, denoted by m-SPRT [16], [22], is based on the principle that one can reject  $H_i$  if it is unlikely to be the answer compared with the current best candidate  $H_{j^*}$ . We stop the test and decide  $H_i$  if  $H_i$  is the only hypothesis left. Let  $H_{j^*}$  denotes the hypothesis that gives the maximum likelihood values of  $X_1, X_2, \dots, X_n$ . m-SPRT constructs binary SPRT  $(a_i, \infty, P_i/P_{j^*})$  for  $i \neq j^*$  such that we reject  $H_i$  if

$$\prod_{k=1}^n \frac{P_i(X_k)}{P_{j^*}(X_k)} < a_i.$$

The error probability  $\epsilon_i$  of rejecting  $H_i$  can also be specified by  $\{a_1, a_2, \dots, a_m\}$  such that

$$\epsilon_i \leq \sum_{j \neq i} a_j.$$

Comparing this approach with the previous one, we see that m-SPRT has substantial saving on computation, since the number of hypotheses decreases once some hypothesis has been rejected. On the other hand, the number of hypothesis for the positive approach remains the same all time. For this reason, we choose m-SPRT in our experiments.

### Maximum-likelihood Alternative

The multihypothesis test can be constructed by using the following binary composite hypothesis test:

$$H : H_j \text{ is true} \quad \text{v.s.} \quad K : \text{One of others is true.}$$

The test is called “composite” because the alternative hypothesis  $K$  has more than one candidate. The likelihood ratio of a composite test can be generalized from a simple-to-simple test by modifying the likelihood function of the alternative hypothesis. For example, the positive approach stops the test and accepts  $H_j$  whenever

$$\frac{\prod_{k=1}^n P_j(X_k)}{\max_{i \neq j} \prod_{k=1}^n P_i(X_k)} > b_j,$$

where  $X_1, X_2, \dots, X_n$  are i.i.d observed samples. On the other hand, the negative approach rejects  $H_j$  once

$$\frac{\prod_{k=1}^n P_j(X_k)}{\max_{i \neq j} \prod_{k=1}^n P_i(X_k)} < a_j.$$

This approach is also known as the *generalized likelihood ratio test* [31].

### Geometric Mean Alternative

Another approach known as the Generalized Sequential Probability Ratio Test (GSPRT) [13] was suggested to use the geometric mean of all the likelihood functions with the negative approach. The resulting test statistics

$$\frac{\prod_{k=1}^n P_j(X_k)}{[\prod_{i=1}^m \prod_{k=1}^n P_i(X_k)]^{1/m}}$$

is compared with the decision boundaries

$$\frac{1 - \epsilon_{j,j}}{[\prod_{i=1}^m (1 - \epsilon_{i,j})]^{1/m}},$$

where  $\epsilon_{i,j}$  is the probability of accepting  $H_j$  when  $H_i$  is true.

### Algebraic Mean Alternative

The so-called M-ary sequential probability ratio test (MSPRT) reported in [7] adopts the positive approach and uses the test statistics based on the Bayes rule by assuming known a priori probabilities  $p(H_1), p(H_2), \dots, p(H_m)$ . The test statistics for alternative hypothesis are a posteriori probabilities for all  $H_j$ . We compare

$$\frac{p(H_j) \prod_{k=1}^n P_j(X_k)}{\sum_{i=1}^m p(H_i) \prod_{k=1}^n P_i(X_k)},$$

with the decision boundaries  $1/(1 + A_j)$  for  $1 \leq j \leq m$ . The probability of wrongly accepting  $H_j$  is bounded by

$$\sum_{i \neq j} p(H_i) \epsilon_{i,j} \leq p(H_j) A_j.$$

## 4 Reference Phase Invariant Classifiers

The a posteriori probabilities of received QAM waveforms given in previous sections are parameterized by  $\vec{p}$ , i.e. a vector of communication parameters such as the carrier frequency, carrier phase, etc. In practice,  $\vec{p}$  is unknown and has to be estimated from received data. For the QAM classification problem, research effort has focused on deriving a reference phase free test statistics [20], [24]. The study of how the reference phase effects the test statistics is valuable because an incorrect carrier frequency estimate will result in error in the phase term. This section will concentrate on how a reference phase invariant test statistic is derived, and how it can be modified to be less sensitive to the frequency estimate error. From the perspective of statistics, a test with unwanted parameters is known as a test with nuisance parameters which is denoted by  $\theta$  in the following.

### 4.1 Nuisance-Parameter-Free Test

There are three approaches reported to solve this problem [4]. One approach is to replace the unwanted parameter  $\theta$  by its maximum likelihood estimate

$$\hat{\theta} = \arg \max_{\theta} p_i(x_1, x_2, \dots, x_k; \theta) \quad (9)$$

and

$$H_i : p_i(x_1, x_2, \dots, x_k; \hat{\theta}), \quad \text{for } i = 1, 2, \dots, m, \quad (10)$$

where  $p_i(x_1, x_2, \dots, x_k; \theta)$  is the probability density function parameterized by a nuisance parameter  $\theta$  if  $H_i$  is true.

Wald suggested to use the average probability density function by introducing a weighting function to average out the unwanted parameter, i.e.

$$H_i : \bar{p}_i(x) = \int_{\theta} p_i(x; \theta) w_i(\theta) d\theta, \quad \text{for } i = 1, 2, \dots, m,$$

where

$$\int_{\theta} w_i(\theta) d\theta = 1.$$

Usually,  $w_i(\theta)$  is chosen to be equally weighted if there is no prior knowledge of the unwanted parameter  $\theta$ . For example, we may assign  $w(\theta_c) = 1/2\pi$ , where  $\theta_c$  is the unknown reference phase of the received signal. In particular, if the average of reference phase is taken by one symbol period, the phase information is lost and the resulting probability density represents the amplitude distribution. Therefore, the averaging process has to be taken for a several-symbol period in order to keep the phase information, i.e.

$$H_i : \bar{p}_i(x_1, x_2, \dots, x_k) = \int_{\theta} p_i(x_1, x_2, \dots, x_k; \theta) w_i(\theta) d\theta. \quad (11)$$

The third approach is to transform the sample sequence  $x_1, x_2, \dots$  to another sequence  $y_1, y_2, \dots$  so that the resulting likelihood function of the new sequence is not parameterized by the nuisance parameter. For example, the phase difference classifier for MPSK signals discussed in [21] uses the transformation

$$\Delta\theta_i = \theta_i - \theta_{i-1} \pmod{2\pi}$$

to obtain the phase difference sequence  $\Delta\theta_i$ , where  $\theta_i$  is the phase of the received signal sampled at the  $i$ th symbol period.

## 4.2 Practical Classifiers Based on Windowed Data

In practice, the carrier frequency might be unknown and has to be estimated from the received signals. The frequency offset due to the estimation error causes the rotation of signal constellation so that the constant reference phase assumption can be violated. Depending on the degree of the frequency estimate error, we can still assume a constant reference phase for a period of time, e.g.  $k$ -symbol period. The immunity to frequency estimate error is determined by the degree that the constant reference assumption is violated in classifier design. The proposed classifiers segment the received waveform into several windows. Data within each window are used to calculate the ML phase estimate by using (9) and/or the likelihood for each hypothesis by (10) or (11). The final test statistic is the product of those test statistics calculated for each set of windowed data. A smaller window size gives a lower correct classification rate since the test statistic contains more reference phase uncertainty. However, it gives a slower degradation of the correct rate in the presence of larger frequency estimate error. The new algorithms based on the windowed data are summarized as follows.

### Algorithm 1 Generalized maximum-likelihood approach

1. Reset the corresponding accumulating score for hypothesis  $H_i$ , i.e.  $L_i = 0, \forall i$ .
2. Collect  $k$  samples  $x_1, x_2, \dots, x_k$ .
3. Calculate the likelihood of hypothesis  $H_i : p_i^*(x_1, x_2, \dots, x_k), \forall i$  by Eq. (10).
4. Update  $L_i$  by  $L_i + \log p_i^*(x_1, x_2, \dots, x_k), \forall i$ .
5. Compute the stopping criterion according to the stopping rules discussed in Section 3. If the stopping criterion is not satisfied, go to Step 2; otherwise, go to the Step 6.
6. Make decision according to the decision rules presented in Section 3.

**Algorithm 2 Maximum average-likelihood approach**

1. Reset the corresponding accumulating score for hypothesis  $H_i$ , i.e.  $L_i = 0, \forall i$ .
2. Collect  $k$  samples  $x_1, x_2, \dots, x_k$ .
3. Calculate the likelihood of hypothesis  $H_i : \bar{p}_i(x_1, x_2, \dots, x_k), \forall i$  by Eq. (11).
4. Update  $L_i$  by  $L_i + \log p_i^*(x_1, x_2, \dots, x_k), \forall i$ .
5. Compute the stopping criterion according to the stopping rules discussed in Section 3. If the stopping criterion is not satisfied, go to Step 1; otherwise, go to the Step 6.
6. Make decision according to the decision rules presented in Section 3.

The above two algorithms have three advantages: (1) a sequential test is feasible; (2) the window size is adjustable to improve the performance for various channel conditions; and (3) a multihypothesis test is feasible.

## 5 Experimental Results

### Example 1: 8-PSK/16-PSK classification

We show in Fig. 4 the required average sample size (ASN) of SPRT compared with FSST (MAP) to classify 8-PSK/16-PSK for symbol SNR ranging from 8 to 17 dB. Their constellations are shown in Figs. 1 (c) and (d). The desired performance is to achieve 99% individual correct rate when the input modulation schemes are either 8-PSK or 16-PSK for SPRT and 99% average correct rate for



FSST. We see that the efficiency of SPRT is about 3 dB for all SNR values. This means that we only need about one half of the samples on the average to make decision to reach the same correct level with SPRT rather than FSST.

**Example 2: 8-PSK/V.29 (7200bps)/Star 8-QAM classification**

The test set is composed of three 8-QAM constellations: 8-PSK, V.29 (7200 bps) and Star 8-QAM. The corresponding constellation diagrams are given in Figs. 1 (c), (d) and (f). Two thousands sets of data were simulated for every constellation. The average and individual correct rates for using 100 symbols for classification is shown in Fig. 5. We can see from the figure that the individual error rates are not evenly distributed. The low SNR correct rate is very low for V.29 but high for 8-PSK in comparison with the average correct rate. The reason why V.29 is difficult to identify can be explained by the fact that V.29 has two amplitude levels, which is the same as Star 8-QAM, and 8 phase angles, which is the same as 8-PSK. As a result, V.29 has a greater chance to be misclassified. This experiment represents a typical challenge to multihypothesis FSST. This problem could be resolved by adjusting the decision threshold of test statistics so that errors can be more evenly distributed.

To demonstrate the performance of SPRT, we generate 1000 sequences of a maximum length of 50,000 symbols for each of the three modulation schemes. By enforcing the maximum number of samples used to make decisions to be finite, we basically adopt truncated SPRT. Furthermore, the following three cases using the negative approach with the maximum-likelihood alternative are examined:

- Test Case 1: symbol SNR 4 dB, known reference phase,  $a_P = a_V = a_S = 0.01$ ;
- Test Case 2: symbol SNR 4 dB, known reference phase,  $a_P = a_V = a_S = 0.05$ ;
- Test Case 3: symbol SNR 0 dB, known reference phase,  $a_P = a_V = a_S = 0.01$ .

Note that the subscripts  $P$ ,  $V$  and  $H$  denote the three modulation schemes 8-PSK, V.29 (7200 bps) and Star 8-QAM, respectively. Also, the choice of boundary decision parameter  $a_P = a$  implies a correct classification rate close to  $1 - 2a$  when the input signal is 8-PSK modulated. Thus, without truncation, the correct rates for Test Cases 1 and 3 should be about 98% while the correct rate for Test Case 2 is about 90%.

Results for Test Case 1 are summarized in Table 1, where RR and AST represent the rejection rate and the average stopping time, respectively. For example, the last 2 columns in Table 1(a) tells

that, with 1,000 Star 8-QAM trial sequences, it takes on the average 120 samples (symbols) to reject  $H_P$  (8-PSK) and the rejection rate is 99.8%, and takes on the average 1,448 samples to reject  $H_V$  (V.29) and the rejection rate is 98.6%. Finally, only 1.6% of the trials make mistakes by rejecting the underlying  $H_S$  while the average stopping time is 961 samples. Note that to make the final decision among 3 hypotheses, we have to reject two hypotheses. The average sample size required to make the final decision is listed at the bottom row for each input signal type. It is clear that we need six times more samples to recognize Star 8-QAM and V.29 than 8-PSK. This is consistent with our previous experience in FSST that 8-PSK is the most recognizable constellation among the three test modulations. The predicted average reject time by using Wald's approximations is shown in Table 1(b) for comparison. The predictions are very close. Table 1(c) shows the decision results of using truncated SPRT. The "correct rate" column shows the individual average correct classification rate. We observe correct rates around 98% via design. The column  $P + V$  denotes that only Hypothesis  $S$  is rejected after 50,000 samples and  $P + V + S$  indicates that no hypothesis has been rejected. It turns out that none of these cases exist in the current experiment.

Test Case 2 is used to show how the values of  $a_P$ ,  $a_V$  and  $a_S$  control the desired performance. By changing their values from 0.01 to 0.05, we see from Table 2 that the correct rate is reduced from 98% to 90%. However, the average sample size is also reduced by a factor of 60 – 75%. For comparison, we also performed FSST which is comparable with Test Case 2 by fixing the sample size to be 900 symbols. The average correct rate is 91.6% and the individual correct rates for 8-PSK, V.29 and Star 8-QAM are 99.9%, 88.0% and 86.8%, respectively, as shown in the last column in Table 2(c). The average correct rate of FSST is slightly lower than that of SPRT (92.9%). The average ASN for FSST and SPRT are 900 and 626, respectively. We see that on the average SPRT needs only about 2/3 of symbols required by FSST with almost identical average correct rates in this experiment.

Results for Test Case 3 are given in Table 3. The average stopping time conditioned on a particular hypothesis and the average sample size to make decision increase significantly due to the lower symbol SNR value. Furthermore, there exists more than one non-rejected hypotheses after 50,000 samples. For example, there are 4.6% of the V.29 simulation sequences which cannot be distinguished from Star 8-QAM. In computing the final correct rate, we are forced to make decision by comparing their probabilities based on 50,000 samples. It turns out the final correct rate is only slightly less the sum of the actual correct rate and the probability of the ambiguity case. For

example, as indicated in the row for V.29 of Table 3 (c), the final correct rate is 97.5% which is approximately equal to the sum of 93.5% (the correct rate) and 4.6% (ambiguity between V+S). Thus, a direct probability comparison based on 50,000 samples is in favor of the right decision. The same statement applies to Star 8-QAM test sequences. This is consistent with our intuition.

**Example 3: Reference phase invariant SPRT**

This experiment includes 11 QAM modulation schemes: BPSK(2P), QPSK(4P), 8-PSK(8P), 16-PSK(16P), V.29, 7200 bps (8V), V.29, 9600 bps(16V), 16-QAM(16Q), 32-QAM(32Q), 64-QAM(64Q), 128-QAM(128Q) and 256-QAM(256Q), where names in the parentheses are their simplified notations. One thousand symbol sequences are generated for every modulation scheme at 10 dB symbol SNR. Results of using SPRT to achieve 99% correct rejection rate with the maximum average-likelihood approach over a 100 symbol window are given in Table 4. Read's decision boundaries are used by applying SPRT after receiving 100 symbols. Therefore, the minimum stopping time is 100 symbol periods. The rejection rate (RR) for every hypothesis and modulation source, the average rejection time (ART) by simulation, and the average correct rate (ACR) and the decision confusion matrix are shown in Table 4(a)-(c), respectively.

**6 Conclusions**

Several QAM classification algorithms based on the likelihood statistics were examined in this research. In particular, we proposed a new QAM classification algorithm based on the sequential probability ratio test (SPRT) and demonstrated that SPRT has several advantages over the classical fixed sample size test (FSST). FSST guarantees the minimum total error probability solution by choosing the hypothesis which gives the maximum a posteriori probability (MAP). However, it does not have the control of the individual error probability given that one of the hypotheses is true. An example was given in the classification of three 8-QAM modulation schemes (8-PSK, V.29 and Star), which showed a typical difficulty of multihypothesis tests, i.e. unbalanced error probability. In contrast, SPRT provides an effective way to control the individual error probability. The number of samples required to make decision represents the delay to establish communication links. For economic reasons, it is desirable to use less amount of data to make decision. SPRT guarantees the minimum average delay with the same level of error probability. The computational complexity is proportional to the number of samples used to make decision, and SPRT also gives the least

amount of computation on the average with the same level of error probability.

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hypothesis	input signals					
	8-PSK		V.29, 7200 bps		Star 8-QAM	
	RR (%)	ART	RR (%)	ART	RR (%)	ART
8-PSK	0.6	199	99.2	207	99.8	120
V.29,7200 bps	99.4	229	1.4	1,066	98.6	1,448
Star 8-QAM	100.0	122	99.4	1,435	1.6	961
ASN	231		1465		1445	

(a)

hypothesis	input signals		
	8-PSK	V.29, 7200 bps	Star 8-QAM
	predicted ART	predicted ART	predicted ART
8-PSK	NA	224	118
V.29,7200 bps	231	NA	1,484
Star 8-QAM	123	1,493	NA

(b)

input signals	CR	decisions						
		P	V	S	P+V	P+S	V+S	P+V+S
8-PSK (P)	99.4	99.4	0.6	0.0	0.0	0.0	0.0	0.0
V.29, 7200 bps (V)	98.6	0.8	98.6	0.6	0.0	0.0	0.0	0.0
Star 8-QAM (S)	98.4	0.2	1.4	98.4	0.0	0.0	0.0	0.0

(c)

Table 1: (a) Rejection rates (RR), average rejection time (ART) and average sample sizes (ASN), (b) predicted average rejection time (ART) and (c) decision results and correct rates (CR) for Test Case 1.

hypothesis	input signals					
	8-PSK		V.29, 7200 bps		Star 8-QAM	
	RR (%)	ART	RR (%)	ART	RR (%)	ART
8-PSK	5.6	114	96.8	131	99.7	81
V.29,7200 bps	94.4	148	8.0	554	92.5	916
Star 8-QAM	100.0	88	95.2	789	7.8	619
ASN	157		820		901	

(a)

hypothesis	input signals		
	8-PSK	V.29, 7200 bps	Star 8-QAM
	predicted ART	predicted ART	predicted ART
8-PSK	NA	146	80
V.29,7200 bps	151	NA	965
Star 8-QAM	77	971	NA

(b)

input signals	SPRT								FSST
	CR	P	V	S	P+V	P+S	V+S	P+V+S	CR
8-PSK (P)	94.4	94.4	5.6	0.0	0.0	0.0	0.0	0.0	99.9
V.29, 7200 bps (V)	92.0	3.2	92.0	4.8	0.0	0.0	0.0	0.0	88.0
Star 8-QAM (S)	92.2	0.3	7.5	92.2	0.0	0.0	0.0	0.0	86.8

(c)

Table 2: (a) Rejection rates (RR), average rejection time (ART) and average sample sizes (ASN), (b) predicted average rejection time (ART) and (c) decision results and correct rates (CR) for Test Case 2.

hypothesis	input signals					
	8-PSK		V.29, 7200 bps		Star 8-QAM	
	RR (%)	ART	RR (%)	ART	RR (%)	ART
8-PSK	1.4	2,714	99.1	2,785	100.0	1,545
V.29, 7200 bps	98.6	2,952	1.9	11,311	94.0	19,541
Star 8-QAM	100.0	1,736	94.4	18,393	2.3	7,496
ASN	2,996		20,218		20,429	

(a)

hypothesis	input signals		
	8-PSK	V.29, 7200 bps	Star 8-QAM
	predicted ART	predicted ART	predicted ART
8-PSK	NA	2,979	1,609
V.29,7200 bps	3,068	NA	21,477
Star 8-QAM	1,675	21,650	NA

(b)

input signals	CR	decisions						
		P	V	S	P+V	P+S	V+S	P+V+S
8-PSK (P)	98.6	98.6	1.4	0.0	0.0	0.0	0.0	0.0
V.29, 7200 bps (V)	97.5	0.9	93.5	1.0	0.0	0.0	4.6	0.0
Star 8-QAM (S)	97.5	0.0	2.3	94.0	0.0	0.0	3.7	0.0

(c)

Table 3: (a) Rejection rates (RR), average rejection time (ART) and average sample sizes (ASN), (b) predicted average rejection time (ART) and (c) decision results and correct rates (CR) for Test Case 3.



source	hypotheses RR (%)										
	2P	4P	8P	16P	8V	16V	16Q	32Q	64Q	128Q	256Q
2P	0.0	100.0	100.0	100.0	100.0	100.0	100.0	100.0	100.0	100.0	100.0
4P	100.0	0.0	100.0	100.0	100.0	100.0	100.0	100.0	100.0	100.0	100.0
8P	100.0	100.0	1.4	98.6	100.0	100.0	100.0	100.0	100.0	100.0	100.0
16P	100.0	100.0	99.2	0.8	100.0	100.0	100.0	100.0	100.0	100.0	100.0
8V	100.0	100.0	100.0	100.0	0.4	100.0	99.6	100.0	100.0	100.0	100.0
16V	100.0	100.0	100.0	100.0	100.0	1.0	100.0	100.0	100.0	100.0	99.0
16Q	100.0	100.0	100.0	100.0	100.0	99.0	4.0	99.0	98.0	100.0	100.0
32Q	100.0	100.0	100.0	100.0	100.0	100.0	100.0	1.4	100.0	99.4	99.2
64Q	100.0	100.0	100.0	100.0	100.0	100.0	99.0	100.0	5.0	98.0	94.0
128Q	100.0	100.0	100.0	100.0	100.0	100.0	100.0	99.0	100.0	2.0	99.0
256Q	100.0	100.0	100.0	100.0	100.0	100.0	100.0	100.0	98.0	98.8	3.2

(a)

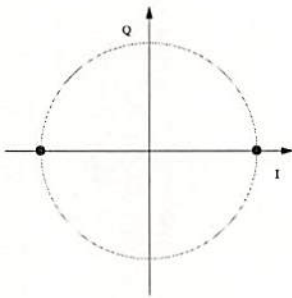
source	ASN	hypotheses ART (simulation)										
		2P	4P	8P	16P	8V	16V	16Q	32Q	64Q	128Q	256Q
2P	100	NA	100	100	100	100	100	100	100	100	100	100
4P	101	100	NA	100	100	101	100	100	100	100	100	100
8P	6,222	100	100	6,375	6,218	100	101	109	113	105	110	105
16P	6,863	100	101	6,852	8,900	100	102	107	112	104	109	104
8V	146	100	100	100	100	100	118	144	105	135	106	132
16V	717	100	100	100	100	110	320	367	375	653	403	582
32Q	9,799	100	100	106	106	101	420	416	11,850	632	9,651	568
64Q	34,385	100	100	103	103	119	556	1,417	615	15,540	540	31,403
128Q	9,312	100	100	107	106	103	450	372	9,298	633	2,550	663
256Q	33,086	100	100	104	104	114	576	976	534	32,970	547	25,267

(b)

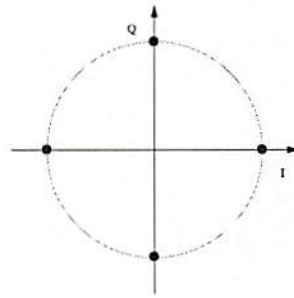
source	ACR	decisions (%)										
		2P	4P	8P	16P	8V	16V	16Q	32Q	64Q	128Q	256Q
2P	100.0	100.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0
4P	100.0	0.0	100.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0
8P	98.6	0.0	0.0	98.6	1.4	0.0	0.0	0.0	0.0	0.0	0.0	0.0
16P	99.2	0.0	0.0	0.8	99.2	0.0	0.0	0.0	0.0	0.0	0.0	0.0
8V	99.6	0.0	0.0	0.0	0.0	99.6	0.0	0.4	0.0	0.0	0.0	0.0
16V	99.0	0.0	0.0	0.0	0.0	0.0	99.0	0.0	0.0	0.0	0.0	1.0
16Q	96.0	0.0	0.0	0.0	0.0	0.0	1.0	96.0	1.0	2.0	0.0	0.0
32Q	98.6	0.0	0.0	0.0	0.0	0.0	0.0	0.0	98.6	0.0	0.6	0.8
64Q	95.0	0.0	0.0	0.0	0.0	0.0	0.0	1.0	0.0	95.0	2.0	2.0
128Q	98.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	1.0	0.0	98.0	1.0
256Q	96.8	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	2.0	1.2	96.8

(c)

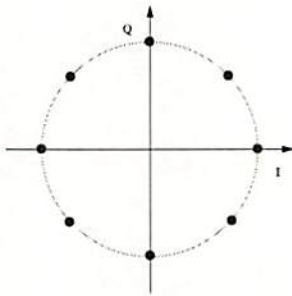
Table 4: (a) Rejection rates (RR), (b) average reject time (ART) and (c) decision results by using reference phase invariant SPRT.



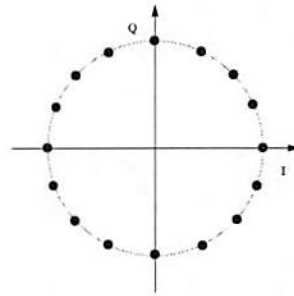
(a) BPSK



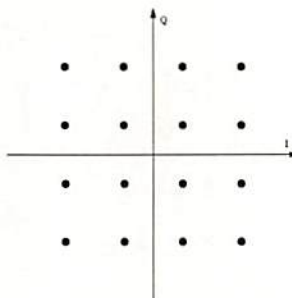
(b) QPSK



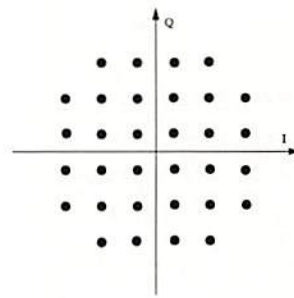
(c) 8-PSK



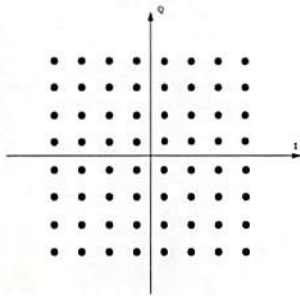
(d) 16-PSK



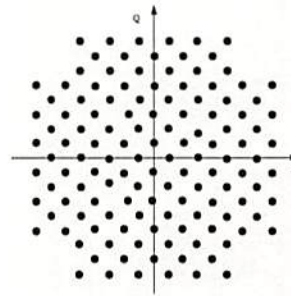
(e) 16-QAM



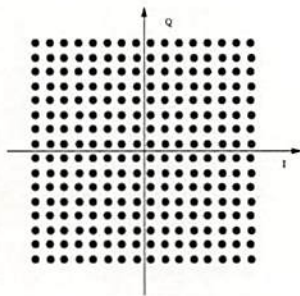
(f) 32-QAM



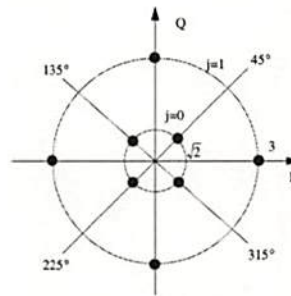
(g) 64-QAM



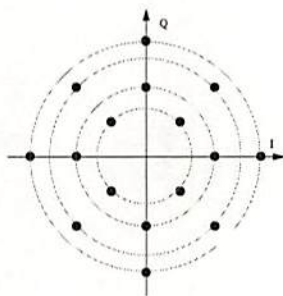
(h) 128-QAM



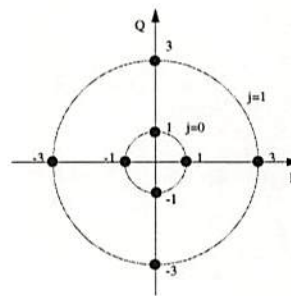
(i) 256-QAM



(j) V29, 7200 bps

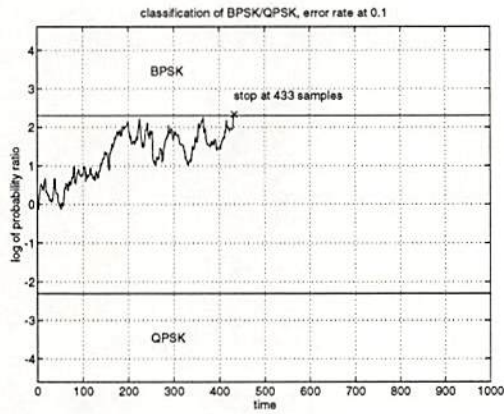


(k) V29, 9600 bps

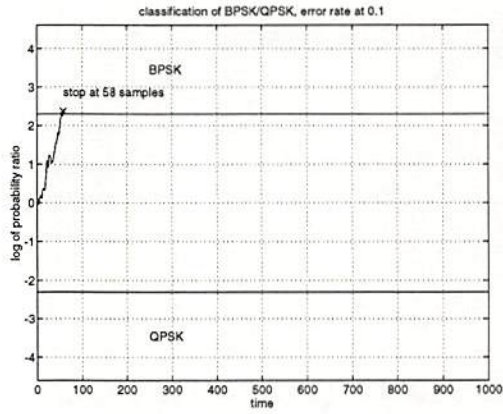


(l) Star 8-QAM

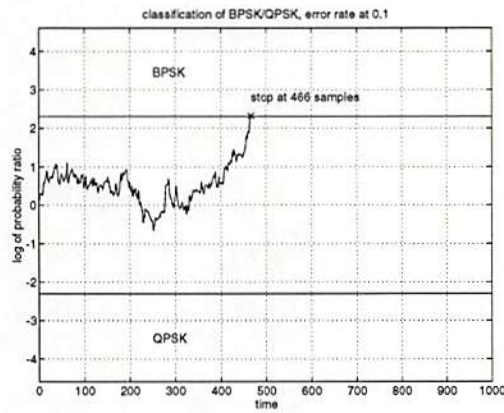
Figure 1: Examples of QAM constellations.



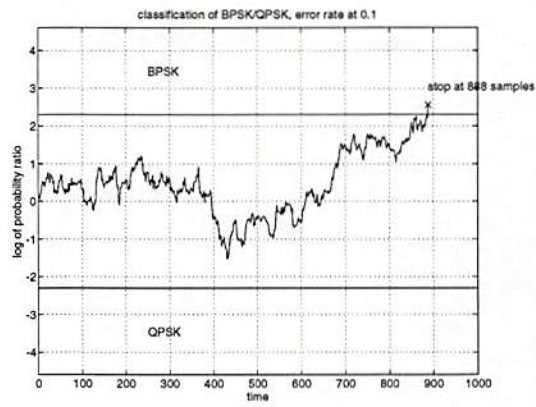
(a) Snapshot 1



(b) Snapshot 2

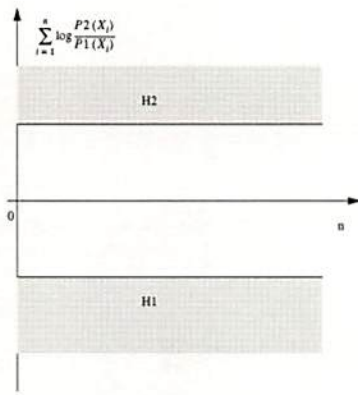


(c) Snapshot 3

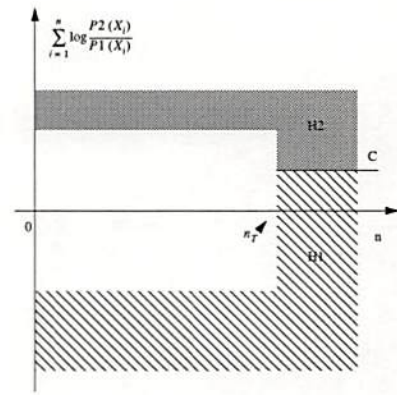


(d) Snapshot 4

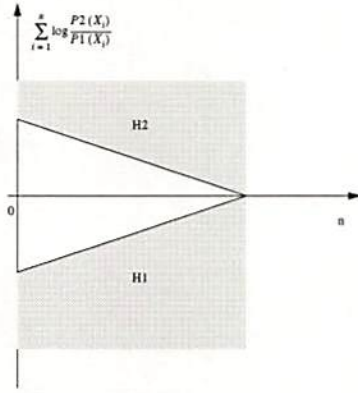
Figure 2: Four snapshots of SPRT for BPSK/QPSK classification with symbol SNR = -7dB.



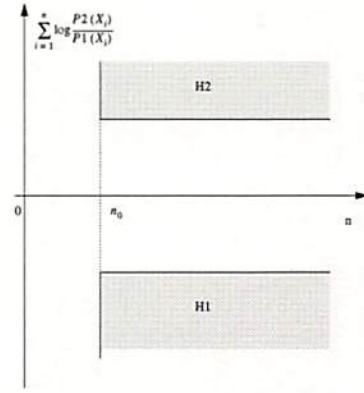
(a)



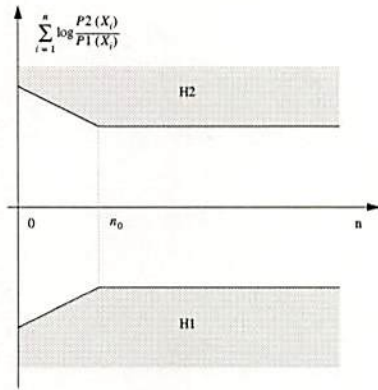
(b)



(c)

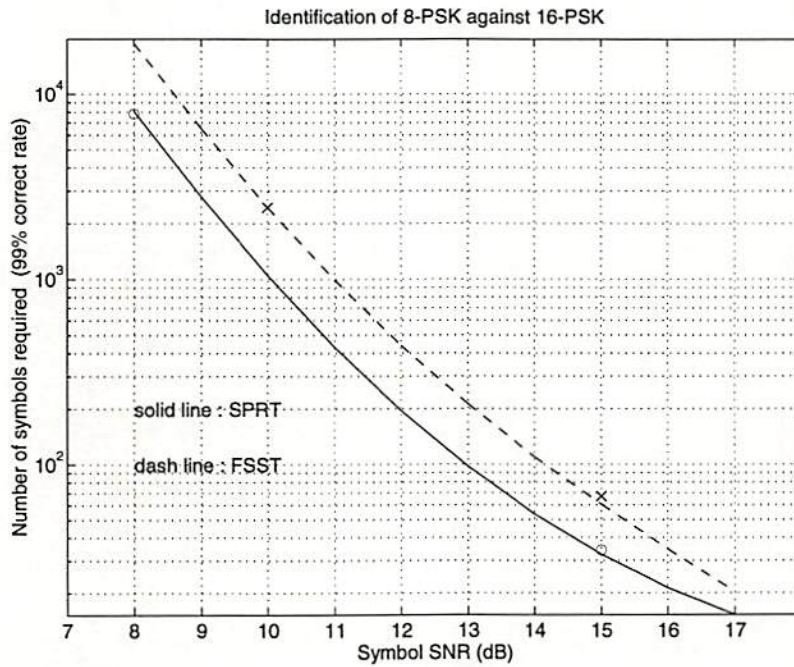


(d)

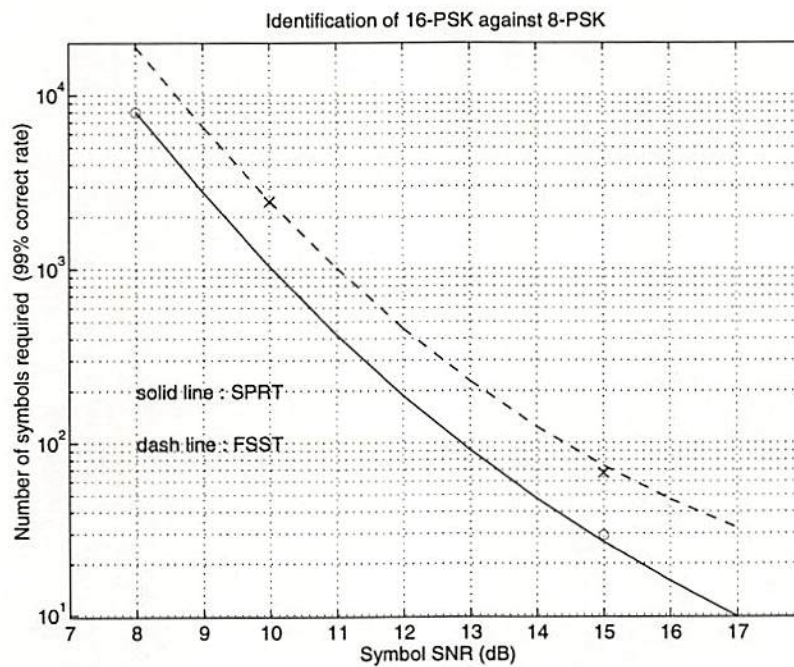


(e)

Figure 3: Examples of decision boundaries: (a) Wald's decision boundary, (b) decision boundary of truncated SPRT, (c) converging decision boundary, (d) Read's decision boundary, and (e) Baruah-Bhattacharjee's decision boundary.



(a)



(b)

Figure 4: The number of symbols required to achieve error level of 0.01 is plotted as a function of symbol SNR in 8-PSK/16-PSK classification with the input is (a) 8-PSK and (b) 16-PSK modulated.

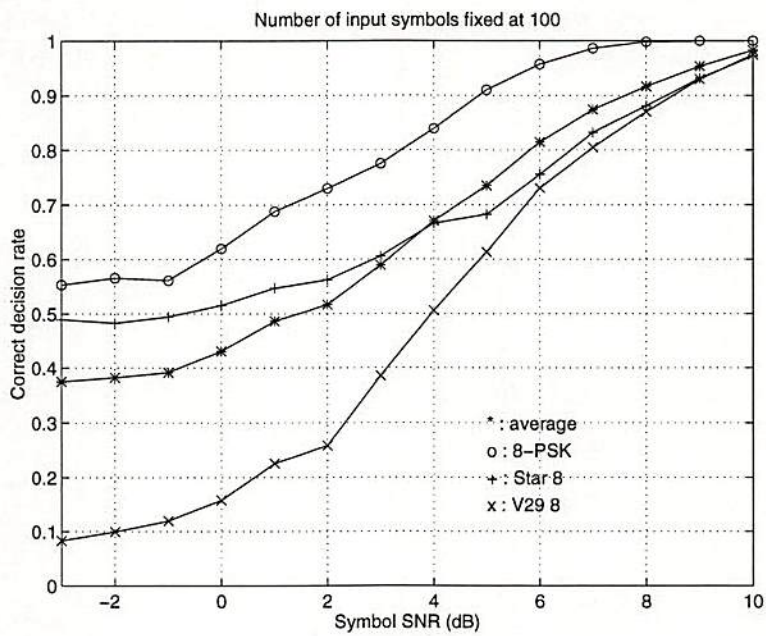


Figure 5: Correct classification rates in classifying 8-PSK, V.29 (7200 bps) and Star 8-QAM using FSST.