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# Optimizations of Multi-hop Cooperative Molecular Communication in Cylindrical Anomalous-Diffusive Channel

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## Abstract

In this paper, the optimizations of multi-hop cooperative molecular communication (CMC) system in cylindrical anomalous-diffusive channel in three-dimensional environment are investigated. First, we derive the performance of bit error probability (BEP) of CMC system under decode-and-forward relay strategy. Then for achieving minimum average BEP, the optimization variables are detection thresholds at cooperative nodes and destination node, and the corresponding optimization problem is formulated. Furthermore, we use conjugate gradient (CG) algorithm to solve this optimization problem to search optimal detection thresholds. The numerical results show the optimal detection thresholds can be obtained by CG algorithm, which has good convergence behaviors with fewer iterations to achieve minimized average BEP compared with gradient decent algorithm and Bisection method which are used in molecular communication.

*Keywords:* Cooperative molecular communication, Cylindrical anomalous-diffusive channel, Optimizations, Conjugate gradient algorithm

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#### **1. Introduction**

The field of molecular communication (MC) [1-2] attracts more attentions of many researchers because of its application prospects in biological environments and industrial fields [3-5]. Recently, more studies have been focused on MC with diffusion in an unbounded fluid environment [6-7].

However, MC in an unbounded environment has some limitations in the practical applications due to its channel characteristics, such as in the blood vessel of human body [8]. Then MC in bounded environment has attracted a lot of attentions of many researchers. The authors [9-10] analytically derived the concentration Green's function (CGF) of MC system in biological cylindrical channel. Based on the Poiseuille flow and the Robin boundary condition, the authors [11] utilized a Markovian-based channel model for MC system to deduce the probability density function. In 2022, Dhok et al. [12] evaluated the performance of probability error of a cooperative molecular communication (CMC) system in cylindrical environment by using a fusion center.

Usually, the molecules propagate in the channel which follows the conventional Brownian motion. However, the scenario of the anomalous diffusion is more extensive comparing with conventional Brownian motion. In 2017, Mai et al. [13] proposed an algorithm for optimizing the network throughput in MC system with anomalous diffusion. In 2019, Trinh et al. [14] derived the first passage time of the molecules by using timing modulation. In 2020, Chouhan et al. [15] calculated the bit error probability (BEP) based on deriving the expression of first passage time density of MC system. In 2021, Trinh et al. [16] derived the observed molecules and the closed-form expressions of the bit error rate. In 2022, the formulation of the first hitting time density of anomalous-diffusive MC system was deduced in [17]. The authors in [18] considered the anomalous diffusion and derived the expression of the CGF of an underlay cognitive MC system.

As so far, some traditional algorithms are used to obtain optimal decision threshold in order to achieve minimum BEP of MC system. Tavakkoli et al. [19] minimized the BEP of two-hop MC system by using bisection algorithm to obtain optimal detection threshold. Chouhan et al. [20] implemented gradient descent (GD) optimization for finding the solutions of the optimization problem which are the values of optimization variables including optimal decision threshold of MC system. Cheng et al. implemented particle swarm optimization algorithm [21] and adaptive genetic algorithm [22] to solve different optimization problems in order to achieve minimum BEP of MC system.

However, the multi-hop CMC system in cylindrical anomalous-diffusive channel in threedimensional (3D) environment has not been studied. On one hand, because of multiple relay nodes in the CMC system, there are multiple detection thresholds needed to be computed by using the maximum a posteriori (MAP) probability detection method with multiple times. However, all the detection thresholds at cooperative nodes and destination node can be obtained simultaneously by using the optimization algorithm. On the other hand, the detection thresholds are optimized by minimizing the average BEP of the CMC system. Therefore, the optimized detection thresholds can improve the reliability of communication of the CMC system. In order to obtain minimum BEP of this system, how to optimize the detection thresholds at cooperative nodes and destination node is a challenge study. But the existing traditional algorithms needs more iterations. It is important to develop more efficient offline optimization techniques. In this paper, we use the conjugate gradient (CG) algorithm to solve the optimization problem. The main contributions of our paper are concluded as follows:

(1) The decode-and-forward (DF) relaying strategy is implemented at cooperative nodes for

the multi-hop CMC system. Then the formulation of average BEP is derived.

(2) We set up an optimization problem for achieving minimum average BEP with optimization variables which are detection thresholds at cooperative nodes and destination node. Then CG algorithm is adopted to solve this optimization problem for searching the optimal detection thresholds.

(3) The numerical results have revealed that CG algorithm is more efficient in finding the optimal detection thresholds at cooperative nodes and the destination node with fewer iterations compared with GD algorithm and Bisection method.

The rest of our paper is organized as follows. Section 2 presents the CMC system model in 3D environment. In Section 3, the detection thresholds are optimized by CG algorithm. In Section 4, the performances of this CMC system with optimized detection thresholds are evaluated. Section 5 summarizes the paper.

#### 2. The Multi-hop CMC System



Fig. 1. The multi-hop CMC system.

The system model of multi-hop CMC in cylindrical channel with anomalous diffusion and drift in 3D environment is shown in **Fig. 1**. This cylindrical channel with the radius  $\rho_{cy}$  is a semi-infinite cylinder with boundary. This system is composed of one source node S, K cooperative nodes CN<sub>1</sub>, CN<sub>2</sub>, ..., CN<sub>K</sub> and one destination node D, which have fixed locations denoted by  $(\rho_S, \phi_S, z_S)$ ,  $(\rho_{CN_1}, \phi_{CN_1}, z_{CN_1})$ ,  $(\rho_{CN_2}, \phi_{CN_2}, z_{CN_2})$ , ...,  $(\rho_{CN_K}, \phi_{CN_K}, z_{CN_K})$ ,  $(\rho_D, \phi_D, z_D)$ , respectively. Here,  $(\rho, \phi, z)$  describes the radial, azimuthal and axial coordinates of each node, respectively. In addition, we have  $0 \le \rho \le \rho_{cy}$ ,  $0 \le \phi < 2\pi$ . In the axial direction, it extends from origin to infinity, then the range of z is  $0 \le z < \infty$ .

We assume that the transmission time is time-slotted. Each time slot duration is  $T_s$ . The modulation method of ON/OFF keying [23] is adopted. The node S and K cooperative nodes instantaneously release molecules at the beginning of each time slot for transmitting bit 1, while releasing no molecules represents the transmission of bit 0. We also suppose that source node S, K cooperative nodes and node D can keep perfect synchronization in time [24-25].

After the transmission process starts, the DF relaying strategy is adopted at each relay node. Different types of molecules are used in each hop. When time slot *j* begins, the source node S releases molecules with type  $A_1$  for transmitting bit 1. For the DF relaying, when time slot *j* ends, the cooperative node CN<sub>1</sub> receives the transmitted bit, which is forwarded to its adjacent node CN<sub>2</sub> when next time slot begins. Each node CN<sub>k</sub> (*k*=1, 2, ..., *K*) can detect type  $A_k$  molecules from node CN<sub>k-1</sub> at the end of time slot (*j*+*k*-1). Then it releases  $A_{k+1}$  molecule types

to forward the decoded bit to node  $CN_{k+1}$ .  $CN_0$  and  $CN_{K+1}$  represent node S node D, respectively. The cooperative nodes and node D are the passive spherical receivers with the radius  $r_{CN_k}$  (k=1, 2, ..., K) and  $r_D$ , respectively. In each time slot, they can count the number of molecules to make the decision that the received bit is 1 or 0 by comparing with the detection thresholds.

After node S and cooperative nodes release molecules, these molecules diffuse in cylindrical anomalous channel. This process can be modeled based on the type of diffusion phenomenon.  $D_{\alpha}(t)$  represents the instantaneous diffusion coefficient which is computed by [26]

$$D_{a}(t) = \alpha D_{p} t^{\alpha - 1}, \qquad (1)$$

where  $D_p$  represents the diffusion coefficient of molecules.  $\alpha \in [0,2]$  is the diffusion exponent which is defined by the normal diffusion with  $\alpha = 1$ , sub-diffusion with  $\alpha \in [0,1)$  and superdiffusion with  $\alpha \in (1,2]$ .

The CGF under the Robin's boundary condition shows the concentration of molecules at node S at time t with location  $(\rho_S, \phi_S, z_S)$  under given initial time  $t_0$ , which is denoted by  $C_S(t;t_0)$  [9]

$$C_{\rm S}(t;t_0) = \frac{\exp\left(-\frac{(z-z_{\rm S}-v(t-t_0))^2}{4(t-t_0)^{\alpha}D_p} - \xi(t-t_0)\right)}{\sqrt{4\pi(t-t_0)^{\alpha}D_p}}$$
(2)  
 
$$\times \left(\sum_{n=0}^{\infty}\sum_{m=1}^{\infty}Q_{nm}\cos(n(\phi-\phi_{\rm S}))J_n(\lambda_{nm}\rho) \times \exp(-D_p\lambda_{nm}^2(t-t_0)^{\alpha})\delta(t-t_0)\right),$$

where  $z_{\rm S}$  is the coordinate value along z axis. v is the drift velocity.  $\xi$  is the degradation constant.  $\rho_{\rm S}$  and  $\rho_{cy}$  are the radii of node S and the cylindrical channel, respectively.  $t_0$  is the

initial time instant. 
$$Q_{nm} = \frac{L_n J_n(\lambda_{nm} \rho_s)}{N_{nm}}, N_{nm} = \frac{\rho_{cy}}{2} \left( J_n^2(\lambda_{nm} \rho_{cy}) - J_{n-1}(\lambda_{nm} \rho_{cy}) J_{n+1}(\lambda_{nm} \rho_{cy}) \right).$$

 $J_n$  and  $\lambda_{nm}$  are the *n*-th order Bessel function of the first kind and the *m*-th eigenvalues, respectively.  $J_{n-1}$  and  $J_{n+1}$  are defined as  $J_n$ . *n* and *m* are integers.  $L_n$  and  $\delta(t)$  are given as follows:

$$L_{n} = \begin{cases} \frac{1}{\pi}, \ n \ge 1, \\ \frac{1}{2\pi}, \ n = 0, \end{cases}$$
(3)

$$\delta(t - t_0) = \begin{cases} 1, \ t \ge t_0, \\ 0, \ t < t_0. \end{cases}$$
(4)

The probability of the case that one released molecule is observed at node  $CN_1$  at time *t* is obtained by [9]

$$p_{(S,CN_1)}(t;t_0) = \iiint_{V_{CN_1}} C_S(t;t_0) \rho \ d\rho d\phi dz \,.$$
(5)

where  $V_{CN_1}$  is the volume of node  $CN_1$  which a spherical region. Then (5) is approximated as  $p_{(S,CN_1)}(t;t_0) = V_{CN_1}C_S(t;t_0)$ . When  $t_0$  represents the *l*-th time slot in which one molecule emitted by node S, the probability of the case that this one molecule is observed in time slot *j* by node  $CN_1$  is given by

$$p_{(S,CN_1)}^{lj} = V_{CN_1} C_S((j-l)T_s;t_0).$$
(6)

#### 3. Derivation of average BEP of the CMC system

 $N_{\rm S}[l]$  and  $x_{\rm S}^l$   $(1 \le l \le j)$  are used to denote the quantity of released molecules and the transmitted bit by node S in time slot *l*, respectively. In time slot *j*, considering the reception at node CN<sub>1</sub>,  $N_{\rm (S,CN_1)}[j]$  is defined as the quantity of received molecules. Then we have

$$N_{(S,CN_1)}[j] = \sum_{l=1}^{J} N_S[l] x_S^l p_{(S,CN_1)}^{lj} + N_{(S,CN_1)}^{Noise},$$
(7)

where  $N_{(S,CN_1)}^{Noise}$  is the noise generated for this link. It is modelled as a Normal distribution  $\mathcal{N}(0, (\sigma_{(S,CN_1)}^{Noise})^2)$ , which has mean 0 and variance  $(\sigma_{(S,CN_1)}^{Noise})^2$  [25].

When  $N_{\rm S}[l]$  is large enough, according to the central limit theorem [25],  $N_{\rm (S,CN_1)}[j]$  is modelled as Gaussian approximation. Therefore, we have

$$N_{(\mathrm{S},\mathrm{CN}_{1})}[j] \sim \mathcal{N}\Big(\mu_{(\mathrm{S},\mathrm{CN}_{1})}[j], \sigma_{(\mathrm{S},\mathrm{CN}_{1})}^{2}[j]\Big),\tag{8}$$

where  $\mu_{(S,CN_1)}[j]$  is the mean and  $\sigma^2_{(S,CN_1)}[j]$  is the variance of  $N_{(S,CN_1)}[j]$ , which are obtained as follows:

$$\mu_{(S,CN_1)}[j] = \sum_{l=1}^{J} \pi_l N_S[k] p_{(S,CN_1)}^{lj}, \qquad (9)$$

$$\sigma_{(\mathrm{S},\mathrm{CN}_{1})}^{2}[j] = \sum_{l=1}^{J} \left[ \pi_{1} N_{\mathrm{S}}[l] p_{(\mathrm{S},\mathrm{CN}_{1})}^{lj} \left( 1 - p_{(\mathrm{S},\mathrm{CN}_{1})}^{lj} \right) + (N_{\mathrm{S}}[l])^{2} \pi_{1} \pi_{0} (p_{(\mathrm{S},\mathrm{CN}_{1})}^{kj})^{2} \right] + (\sigma_{(\mathrm{S},\mathrm{CN}_{1})}^{\mathrm{Noise}})^{2}, \quad (10)$$

where  $\pi_1$  and  $\pi_0$  are the transmission probabilities of 1 and 0 by node S when each time slot begins, respectively. We have  $Pr(x_s[l]=1) = \pi_1 = 0.5$  and  $Pr(x_s[l]=0) = \pi_0 = 0.5$ .

 $H_0$  and  $H_1$  indicate the cases that when time slot *j* begins, node S transmits bits 0 and 1, respectively. Then the corresponding Normal distributions under  $H_0$  and  $H_1$  are formulated by

$$H_{0}: N_{(S,CN_{1})}[j] \sim \mathcal{N}\left(\mu_{(S,CN_{1})}^{0}[j], (\sigma_{(S,CN_{1})}^{0}[j])^{2}\right),$$

$$H_{1}: N_{(S,CN_{1})}[j] \sim \mathcal{N}\left(\mu_{(S,CN_{1})}^{1}[j], (\sigma_{(S,CN_{1})}^{1}[j])^{2}\right),$$
(11)

where  $\mu_{(S,CN_1)}^w[j]$  (w=0, 1) is the mean and  $(\sigma_{(S,CN_1)}^w[j])^2$  (w=0, 1) is the variance of  $N_{(S,CN_1)}[j]$  under  $H_w$ , respectively. According to (9) and (10),  $\mu_{(S,CN_1)}^w[j]$  and  $(\sigma_{(S,CN_1)}^w[j])^2$  (w=0, 1) can be computed by

$$\mu_{(S,CN_{1})}^{0}[j] = \sum_{l=1}^{j-1} \pi_{1} N_{S}[l] p_{(S,CN_{1})}^{l(j-1)},$$

$$(\sigma_{(S,CN_{1})}^{0}[j])^{2} = \sum_{l=1}^{j-1} \left[ \pi_{1} N_{S}[l] p_{(S,CN_{1})}^{lj} \left( 1 - p_{(S,CN_{1})}^{lj} \right) + (N_{S}[l])^{2} \pi_{1} \pi_{0} (p_{(S,CN_{1})}^{lj})^{2} \right] + (\sigma_{(S,CN_{1})}^{Noise})^{2},$$

$$(12)$$

$$\mu_{(S,CN_{1})}^{1}[j] = N_{S}[j] p_{(S,CN_{1})}^{lj} + \mu_{(S,CN_{1})}^{0}[j],$$

$$(\sigma_{(S,CN_{1})}^{1}[j])^{2} = N_{S}[j] p_{(S,CN_{1})}^{lj} \left( 1 - p_{(S,CN_{1})}^{lj} \right) + (\sigma_{(S,CN_{1})}^{0}[j])^{2}.$$

The detection threshold at node  $CN_1$  is  $\eta_{CN_1}$ . Then when each time slot ends, the detection rule at node  $CN_1$  is written as

$$\hat{x}_{CN_{1}}[j] = \begin{cases} 1, & \text{if } N_{(S,CN_{1})}[j] \ge \eta_{CN_{1}}, \\ 0, & \text{if } N_{(S,CN_{1})}[j] < \eta_{CN_{1}}, \end{cases}$$
(13)

where  $\hat{x}_{CN_1}^j$  is denoted by the bit decoded by node  $CN_1$  when time slot *j* ends. When  $x_S^j \neq \hat{x}_{CN_1}^j$ , an error occurs in this time slot.  $Pr(\hat{x}_{CN_1}^j = 0 | x_S^j = 1)$  and  $Pr(\hat{x}_{CN_1}^j = 1 | x_S^j = 0)$  represent the error probabilities of bits transmission of 1 and 0, respectively. Then we have  $Pr(\hat{x}_{CN_1}^j = 0 | x_{CN_1}^j = 1) - Pr(N_1 = 1 | x_{CN_1}^j = 1)$ 

$$\Pr(x_{CN_{1}}^{j} = 0 | x_{S}^{j} = 1) = \Pr(N_{(S,CN_{1})}[j] < \eta_{CN_{1}} | x_{S}^{j} = 1)$$

$$= Q\left(\frac{\eta_{CN_{1}} - \mu_{(S,CN_{1})}^{0}[j]}{\sigma_{(S,CN_{1})}^{0}[j]}\right),$$

$$\Pr(\hat{x}_{CN_{1}}^{j} = 1 | x_{S}^{j} = 0) = \Pr(N_{(S,CN_{1})}[j] \ge \eta_{CN_{1}} | x_{S}^{j} = 0)$$

$$= Q\left(\frac{\eta_{CN_{1}} - \mu_{(S,CN_{1})}^{1}[j]}{\sigma_{(S,CN_{1})}^{1}[j]}\right),$$
(14)
(14)
(15)

where the function Q(x) is denoted by  $Q(x) = \frac{1}{\sqrt{2\pi}} \int_{x}^{\infty} e^{-\frac{u^{2}}{2}} du$ . Based on formulas (14) and (15),

 $Pe_{(S,CN_1)}[j]$  which represents the BEP of one bit transmission in time slot j is written as

$$Pe_{(S,CN_1)}[j] = \pi_1 \Pr(\hat{x}_{CN_1}^j = 0 | x_S^j = 1) + \pi_0 \Pr(\hat{x}_{CN_1}^j = 1 | x_S^j = 0).$$
(16)

Considering the link  $CN_k \rightarrow CN_{k+1}$  in the (j+k)-th time slot,  $N_{(CN_k, CN_{k+1})}[j+k]$  is the number of received molecules by  $CN_{k+1}$ . The binary hypothesis testing problem based on  $N_{(CN_k, CN_{k+1})}[j+k]$  is established at node  $CN_{k+1}$  as follows:

$$H_{0}: N_{(CN_{k}, CN_{k+1})}[j+k] \sim \mathcal{N}\left(\mu_{(CN_{k}, CN_{k+1})}^{0}[j], (\sigma_{(CN_{k}, CN_{k+1})}^{0}[j])^{2}\right),$$

$$H_{1}: N_{(CN_{k}, CN_{k+1})}[j+k] \sim \mathcal{N}\left(\mu_{(CN_{k}, CN_{k+1})}^{1}[j], (\sigma_{(CN_{k}, CN_{k+1})}^{1}[j])^{2}\right),$$
(17)

where  $\mu_{(CN_k,CN_{k+1})}^w[j]$  (w=0, 1) is the mean and  $(\sigma_{(CN_k,CN_{k+1})}^w[j])^2$  (w=0, 1) is the variance of  $N_{(CN_k,CN_{k+1})}[j+k]$  under  $H_w$ , respectively.

Let  $x_{CN_k}^{j+k} = 0$  and  $x_{CN_k}^{j+k} = 1$  represent the bits transmission of 0 and 1 by  $CN_k$  at the time

slot (j+k), respectively.  $\hat{x}_{CN_{k+1}}^{j+k}$  is 1 and 0 which are decoded by node  $CN_{k+1}$  when time slot (j+k) ends, respectively. Thus the corresponding error probabilities for the link  $CN_k \rightarrow CN_{k+1}$  are defined as  $Pr(\hat{x}_{CN_{k+1}}^{j+k} = 1|x_{CN_k}^{j+k} = 0)$  and  $Pr(\hat{x}_{CN_{k+1}}^{j+k} = 0|x_{CN_k}^{j+k} = 1)$ , respectively, which are computed according to (17) as follows:

$$\Pr(\hat{x}_{CN_{k+1}}^{j+k} = 1 | x_{CN_{k}}^{j+k} = 0) = \Pr(N_{(CN_{k}, CN_{k+1})}[j+k] \ge \eta_{CN_{k+1}} | x_{CN_{k}}^{j+k} = 0),$$
(18)

$$\Pr(\hat{x}_{CN_{k+1}}^{j+k} = 0 | x_{CN_{k}}^{j+k} = 1) = \Pr(N_{(CN_{k}, CN_{k+1})}[j+k] < \eta_{CN_{k+1}} | x_{CN_{k}}^{j+k} = 1),$$
(19)

where  $\eta_{CN_{k+1}}$  is detection threshold at node  $CN_{k+1}$ .

After node  $CN_{k+1}$  decodes the received bit, it forwards  $\hat{x}_{CN_{k+1}}^{j+k}$  to the next relay node  $CN_{k+2}$ . The transmitted bit by  $CN_{k+2}$  at the (j+k+1)-th time slot is  $\hat{x}_{CN_{k+1}}^{j+k+1}$ . Assume the BEP of bit 0 or 1 transmission in time slot *j* from S to  $CN_k$  are  $Pe^0_{(S,CN_k)}[j]$  and  $Pe^1_{(S,CN_k)}[j]$ , respectively. Then considering the link  $S \rightarrow CN_{k+1}$ , the BEP of one bit transmission in time slot *j* are computed by

$$Pe^{0}_{(S,CN_{k+1})}[j] = Pe^{0}_{(S,CN_{k})}[j] \times \Pr[\hat{x}^{j+k}_{CN_{k+1}} = 1 | x^{j+k}_{CN_{k}} = 1] + (1 - Pe^{0}_{(S,CN_{k})}[j]) \times \Pr[\hat{x}^{j+k}_{CN_{k+1}} = 1 | x^{j+k}_{CN_{k}} = 0],$$
(20)

$$Pe_{(S,CN_{k+1})}^{l}[j] = Pe_{(S,CN_{k})}^{l}[j] \times \Pr[\hat{x}_{CN_{k+1}}^{j+k} = 0 \mid x_{CN_{k}}^{j+k} = 0] + (1 - Pe_{(S,CN_{k})}^{l}[j]) \times \Pr[\hat{x}_{CN_{k+1}}^{j+k} = 0 \mid x_{CN_{k}}^{j+k} = 1].$$
(21)

When considering the multi-hop CMC system with k=K in (20) and (21), the BEP of one bit in the *j*-th time slot from node S denoted by  $Pe_{(S,D)}[j]$  is computed by

$$Pe_{(S,D)}[j] = \pi_1 Pe_{(S,CN_{K+1})}^1[j] + \pi_0 Pe_{(S,CN_{K+1})}^0[j].$$
<sup>(22)</sup>

## 4. Optimization of detection thresholds of multi-hop CMC system

The optimization problem of detection thresholds of the multi-hop CMC system in cylindrical channel is expressed as follows:

$$\min_{\eta_{\mathrm{CN}_{1}}, \eta_{\mathrm{CN}_{2}}, \dots, \eta_{\mathrm{CN}_{K}}, \eta_{\mathrm{D}}} Pe_{(\mathrm{S},\mathrm{D})}[j],$$
(23)

where  $\eta_{CN_1}, \eta_{CN_2}, ..., \eta_{CN_K}$  are the detection thresholds of nodes CN<sub>1</sub>, CN<sub>2</sub>, ..., CN<sub>K</sub>, respectively.  $\eta_D$  is the detection threshold at destination node D.

 $\eta_u$  is a vector of detection thresholds at the *u*-th iteration which is represented by  $\eta_u = [\eta_{CN_1}^u, \eta_{CN_2}^u, ..., \eta_{CN_K}^u, \eta_D^u]$ .  $\nabla Pe_{(S,D)}[j](\eta_u)$  is the gradient of  $Pe_{(S,D)}[j]$  with  $\eta_u$  which is computed by

$$\nabla Pe_{(S,D)}[j](\eta_u) = \left[\frac{\partial Pe_{(S,D)}[j](\eta_u)}{\partial \eta_{CN_1}}, \frac{\partial Pe_{(S,D)}[j](\eta_u)}{\partial \eta_{CN_2}}, \dots, \frac{\partial Pe_{(S,D)}[j](\eta_u)}{\partial \eta_{CN_K}}, \frac{\partial Pe_{(S,D)}[j](\eta_u)}{\partial \eta_D}\right], \quad (24)$$

where 
$$\frac{\partial Pe_{(\mathrm{S},\mathrm{D})}[j](\eta_u)}{\partial \eta_{\mathrm{CN}_1}}, \frac{\partial Pe_{(\mathrm{S},\mathrm{D})}[j](\eta_u)}{\partial \eta_{\mathrm{CN}_2}}, \dots, \frac{\partial Pe_{(\mathrm{S},\mathrm{D})}[j](\eta_u)}{\partial \eta_{\mathrm{CN}_K}}, \frac{\partial Pe_{(\mathrm{S},\mathrm{D})}[j](\eta_u)}{\partial \eta_{\mathrm{D}}}$$
 are the first

derivative of  $Pe_{(S,D)}[j](\eta_u)$  with respect to  $\eta_{CN_1}, \eta_{CN_2}, ..., \eta_{CN_K}$ ,  $\eta_D$ , respectively. The CG algorithm is described as follows:

| Algorithm 1. CG Algorithm for Optimizing Detection Thresholds in CMC System  |  |  |  |  |
|--|--|--|--|--|
| <b>Input:</b> The maximum number of iterations is $N=50$ . $\varepsilon = 0.0001$ is the accuracy. Randomly                        |  |  |  |  |
| set a starting point $\eta_0 = [\eta_{CN_1}^0, \eta_{CN_2}^0,, \eta_{CN_K}^0, \eta_D^0]$ which is composed of the initial value of |  |  |  |  |
| detection thresholds at nodes $CN_1, CN_2, \dots, CN_K$ and node D.  |  |  |  |  |
| <b>Output:</b> The vector of optimal detection thresholds $\eta^* = [\eta^*_{CN_1}, \eta^*_{CN_2},, \eta^*_{CN_K}, \eta^*_D]$ .    |  |  |  |  |
| 1: <b>for</b> $u = 0$ to $N$ do  |  |  |  |  |
| 2: <b>if</b> $u \% K == 0$   |  |  |  |  |
| 3: Update decent direction: $\Delta(\eta_{u+1}) = -\nabla P e_{(S,D)}[j](\eta_u)$ .  |  |  |  |  |
| 4: else  |  |  |  |  |
| $ = \left( \nabla P e_{(\mathbf{S},\mathbf{D})}[j](\eta_u) \right)^T \nabla P e_{(\mathbf{S},\mathbf{D})}[j](\eta_u) $             |  |  |  |  |
| 5. $p_{u+1} = \frac{1}{(\nabla Pe_{(S,D)}[j](\eta_{u-1}))^T \nabla Pe_{(S,D)}[j](\eta_{u-1})}$ .                                   |  |  |  |  |
| 6: Update decent direction: $\Delta(\eta_{u+1}) = -\nabla P e_{(S,D)}[j](\eta_u) + \beta_{u+1}\Delta(\eta_u)$ .                    |  |  |  |  |
| 7: end if  |  |  |  |  |
| 8: Choose step size $\gamma_u$ for line search.  |  |  |  |  |
| 9: Update $\eta_{u+1} = \eta_u + \gamma_u \Delta(\eta_{u+1})$ , calculate $Pe_{(S,D)}[j](\eta_{u+1})$ .                            |  |  |  |  |
| 10: Until $\left\  Pe_{(S,D)}[j](\eta_{u+1}) - Pe_{(S,D)}[j](\eta_u) \right\  \le \varepsilon$ . Then output the vector which is   |  |  |  |  |
| composed of optimal detection thresholds $\eta^* = [\eta^*_{CN_1}, \eta^*_{CN_2},, \eta^*_{CN_k}, \eta^*_D]$ .                     |  |  |  |  |
| 11: end for  |  |  |  |  |

It is noted that when  $\|\nabla Pe_{(S,D)}[j](\eta_0)\| > \lambda(\lambda = 10^{-6})$  and the corresponding  $Pe_{(S,D)}[j](\eta_0)$  at  $\eta_0$  is smaller than 0.5, this algorithm can converge fast, and it performs exceptionally well. Then we can get optimized results of detection thresholds at cooperative nodes and destination node. When  $\|\nabla Pe_{(S,D)}[j](\eta_0)\| \le \lambda$  and the corresponding  $Pe_{(S,D)}[j](\eta_0)$  at  $\eta_0$  is equal to 0.5, this algorithm is difficult to achieve convergence. Under such a case, this algorithm will face challenges. In order to solve this problem, we search for a better  $\eta_0$  at the neighborhood of  $\eta_0$  which can satisfy the condition that  $\|\nabla Pe_{(S,D)}[j](\eta_0)\| > \lambda$ . Then  $\eta_0$  is updated as  $\eta_0 = \eta_0$ .

#### 5. Numerical results

The numerical results are shown in this section and the default numerical parameters are given in **Table 1**.

| Parameter  | Value        | Parameter  | Value   |
|--|--------------|------------|---|
| $r_{{ m CN}_k}$ , $r_{ m D}$   | 2µm          | $D_p$      | $1 \times 10^{-10} \text{ m}^2/\text{s}$ [18] |
| $(\rho_{\rm S},\phi_{\rm S},z_{\rm S})$  | (0,0,0)      | v          | 65µm/s <mark>[18]</mark>                      |
| $(\rho_{\text{CN}_1}, \phi_{\text{CN}_1}, z_{\text{CN}_1})$                      | (2µm,0,5µm)  | νy         | $9s^{-1}[12]$                                 |
| $(\rho_{{}_{\mathrm{CN}_2}}, \phi_{{}_{\mathrm{CN}_2}}, z_{{}_{\mathrm{CN}_2}})$ | (2µm,0,10µm) | $ ho_{cy}$ | 5μm <mark>[12]</mark>                         |
| $(\rho_{\rm D},\phi_{\rm D},z_{\rm D})$  | (2µm,0,15µm) | $T_s$      | 500ms   |
| α  | [0,2]        | N          | 3×10 <sup>4</sup>                             |

Table 1. The numerical parameters



Fig. 2. The average BEP vs  $\eta_{\rm CN_1}$  and  $\eta_{\rm D}$ .

 $\eta_{\rm CN_1}$  and  $\eta_{\rm D}$  are the thresholds at node CN<sub>1</sub> and node *D*, respectively. Fig. 2 shows the average BEP is changing with  $\eta_{\rm CN_1}$  and  $\eta_{\rm D}$  when there is one cooperation node CN<sub>1</sub>. When  $\eta_{\rm CN_1}$  takes the same value, the average BEP decreases first and then increases with the increasing value of  $\eta_{\rm D}$ . The change trend under fixed value of  $\eta_{\rm D}$  is the same as that under fixed value of  $\eta_{\rm CN_1}$ . We can see that it is a convex problem when two detection thresholds with  $\eta_{\rm CN_1}$  and  $\eta_{\rm D}$  are optimized. Therefore, CG algorithm can be adopted to solve this optimization problem efficiently. The value of the lowest point with red dots is represented by (4800,7150, 0.0762), which means that the optimized results are  $\eta_{\rm CN_1}$ =4800,  $\eta_{\rm D}$ =7150 and the corresponding average BEP is 0.0762.



Fig. 3. The comparison results of convergence with CG, GD and Bisection algorithms.

In **Fig. 3**, we give the convergence analysis of CG, GD and Bisection algorithms. Under these three algorithms, when the number of iterations is increasing, the performances of average BEP are both decreasing. Finally, they can achieve convergence. We can see that CG algorithm converges fastest than GD algorithm and Bisection algorithm. We can see that for the same parameter setting, the number of iterations of CG algorithm, GD algorithm and Bisection algorithm are 5, 11 and 7, respectively. Therefore, we use CG algorithm to solve this optimization problem with fewer iterations and obtain optimal detection thresholds and minimum average BEP.



Fig. 4. Convergence analysis under different values of (a)  $\alpha$ ; (b)  $\rho_{cv}$ .

When  $\alpha$  and  $\rho_{cy}$  take different values, the convergence results are shown in Fig. 4(a) and Fig. 4(b), respectively. Here  $\alpha$  is the diffusion exponent which has corresponding ranges. The CG algorithm has good convergence which shows that the average BEP can converge with fewer iterations. In Fig. 4(a), when  $\alpha = 0.5$  and  $\alpha = 1.5$ , the corresponding number of iterations are 6 and 5, respectively. In Fig. 4(b), when  $\rho_{cy} = 5\mu m$  and  $\rho_{cy} = 7\mu m$ , the average BEP achieves convergence within 3 iterations and 5 iterations, respectively. Second, when the values of  $\alpha$  are smaller and the values of  $\rho_{cy}$  are larger, the average BEP are both larger.

This result is explained by the facts: on one hand, smaller values of  $\alpha$  results in slower diffusion. On the other hand, larger values of  $\rho_{cy}$  means the same number of molecules can diffuse in much larger space. These two cases both lead to a decrease of the number of molecules received by cooperative nodes CN<sub>1</sub>, CN<sub>2</sub> and node D, finally there is a decrease in BEP for each link.



Fig. 5. The average BEP vs  $\eta_{\rm CN_1}$  under different values of (a)  $\alpha$  ; (b)  $\rho_{\rm cv}$ .

The average BEP is changing with  $\eta_{CN_1}$  under different values of  $\alpha$  and  $\rho_{cy}$  in Fig. 5. Different ranges in Fig. 5(a) including  $\alpha < 1$ ,  $\alpha = 1$  and  $\alpha > 1$  represent sub-diffusion, normal diffusion and super-diffusion, respectively. We can see that when  $\alpha$  is increasing, the average BEP is found to decrease. This is explained as follows: larger value of  $\alpha$  will accelerate the diffusion of molecules. Then the number of molecules arriving at nodes CN<sub>1</sub>, CN<sub>2</sub> and node D will increase. Finally, this reduces the average BEP. In addition, the optimal thresholds at CN<sub>1</sub>, CN<sub>2</sub> and node D are given at the lowest point with minimum value of average BEP. When  $\alpha = 1.5$ ,  $\eta_{CN_1} = 9100$ ,  $\eta_{CN_2} = 10970$ ,  $\eta_D = 11082$  and the corresponding average BEP is 0.097. In Fig. 5(b),  $\rho_{cy} = \{5\mu m, 7\mu m, 9\mu m\}$ . The performance of average BEP is decreasing with  $\eta_{CN_1}$ . When  $\eta_{CN_1}$  achieves some value, the average BEP achieves its minimum value and then increase. Moreover, the number of received molecules is larger when  $\rho_{cy}$  takes smaller value, and the corresponding detection thresholds are also larger to obtain minimum value of average BEP. For the case  $\rho_{cy} = 5\mu m$ ,  $\eta_{CN_1} = 5100$ ,  $\eta_{CN_2} = 9200$ , and  $\eta_D = 12228$ , the average BEP is larger than those under the cases  $\rho_{cy} = 7\mu m$  and  $\rho_{cy} = 9\mu m$ .



Fig. 6. The average BEP vs  $\eta_{CN_1}$  under different distances along z-axial direction for each hop.

We use  $d_0$  to represent the distance between two adjacent nodes along *z*-axial direction. Then consider three-hop CMC system with two cooperation nodes CN<sub>1</sub> and CN<sub>2</sub>, the transmission distance is  $3d_0$ . Fig. 6 gives the result the average BEP is changing with detection thresholds when  $d_0$  takes different values. The change trend of average BEP is similar as in Fig. 5. When  $d_0$  is with larger value which results in lower receiving probability, the average BEP is larger and the detection thresholds at lowest point of average BEP are smaller. When  $d_0=5\mu m$ , the average BEP is 0.386 which is larger than that when  $d_0=3\mu m$ , and the corresponding values of detection thresholds at lowest point of BEP are  $\eta_{CN_1}=3300$ ,  $\eta_{CN_2}=3300$ ,  $\eta_{CN_2}=3300$ ,  $\eta_{CN_2}=3300$ ,  $\eta_{CN_2}=300$ .

=3707,  $\eta_{\rm D}$  =3721 which are smaller than those when  $d_0$ =3 $\mu$ m.

In order to show the differences between DF relay strategy and amplify-and-forward (AF) relay strategy, we give the comparison results in **Fig. 7**. The parameters are set the same for the two relay strategies. Especially, for the AF relay strategy, the amplification factor is set as 5. According to **Fig. 7**, we can see that the average BEP under DF relay strategy decreases faster and is smaller than that under AF relay strategy for the same value of  $\eta_{CN_1}$ . In particular, the detection thresholds at lowest point are the same for these two relay strategies. Therefore, for the CMC system, we choose DF relay strategy.



Fig. 7. The average BEP vs  $\eta_{\rm CN_1}$  with DF and AF relay strategies.



**Fig. 8.** The average BEP versus *K* under different distances between nodes S and D along *z*-axial direction.

 $z_{(S,D)}$  is used to represent the distance between nodes S and D along *z*-axial direction. In **Fig. 8**, the average BEP is decreasing with the value of *K* which is the number of cooperative nodes. When the total distance along *z*-axial direction  $z_{(S,D)}$  is fixed, the distance along *z*-axial direction for each hop is decreasing with increasing value of *K*. The receiving probabilities at each cooperative nodes CN<sub>1</sub>, CN<sub>2</sub> and node D also increase. Finally, the average BEP is decreasing. For each same value of *K*, the average BEP of this CMC system under  $z_{(S,D)} = 20\mu$ m is the lowest than those under  $z_{(S,D)} = 10\mu$ m and  $z_{(S,D)} = 15\mu$ m.

## 6. Conclusion

This paper studied the optimizations of multi-hop CMC system in cylindrical anomalousdiffusive channel in a 3D environment. We have derived the average BEP which is optimization objective function. Then the optimization problem for minimizing the average BEP was effectively solved by CG algorithm. The numerical results have shown that CG algorithm can converge by using fewer iterations in contrast to GD algorithm and Bisection algorithm. In addition, we also have shown that some main parameters have impacts on the optimization results. The lager value of  $\alpha$ , smaller value of  $\rho_{cy}$  and initial distance between two adjacent nodes along z-axial direction, then the smaller value of average BEP. In future work, we intend to explore other traditional and efficient optimization methods for solving the optimization problem under the scenario with mobile nodes in CMC system. In additon, we plan to use data driven methods, such as the deep learning methods to optimize optimal detection thresholds under different system parameters for CMC system.

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