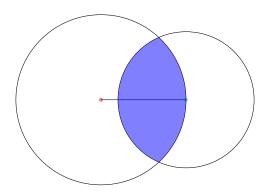
Sojourn Time of Moving Relays in Dual-Hop Cooperative Communication

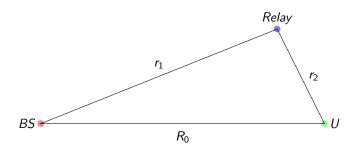
Prudhvi Porandla 110070039

June 20, 2016

Problem



Downlink Cooperation policy



$$\{P_b r_1^{-\gamma} > P_b R_0^{-\gamma}, P_r r_2^{-\gamma} > P_b R_0^{-\gamma}\}$$

 P_b and P_r are the transmission powers of base station and relay respectively.

$$\{r_1 < R_0, r_2 < R_2\}$$
 where $R_2 = cR_0, c = \left(rac{P_r}{P_b}
ight)^{rac{1}{\gamma}}$

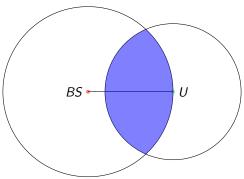


Figure: Feasible region

Motivation

► Relay selection

Motivation

► Relay selection

► Effect on the whole network

Mobility Model

The *n*th transition of a node is defined by the quadruple

$$(\mathbf{X}_{n-1},\mathbf{X}_n,V_n,S_n)$$

 X_{n-1} and X_n are the starting and destination waypoints.

Velocity - V_n , Pause time or Thinking time at destination - S_n

Mobility Model

The *n*th transition of a node is defined by the quadruple

$$(\mathbf{X}_{n-1},\mathbf{X}_n,V_n,S_n)$$

 X_{n-1} and X_n are the starting and destination waypoints.

Velocity - V_n , Pause time or Thinking time at destination - S_n

Mobility model

- ▶ Distribution of transition length($L_n = \|\mathbf{X}_{n-1} \mathbf{X}_n\|$)
- ▶ Distribution of angle made by the vector $\mathbf{X}_n \mathbf{X}_{n-1}$ w.r.t x-axis.
- ▶ Distributions of V_n and S_n

Rayleigh RWP

Angle is chosen uniformly from $[0,2\pi]$

Transition length is Rayleigh distributed with parameter λ .

$$P(L>I) = exp(-\lambda \pi I^2), I \ge 0$$

$$V \equiv v$$
 and $S \equiv 0$

Rayleigh RWP

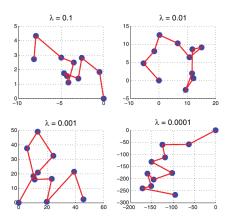


Figure: Sample traces of Rayleigh RWP¹



¹Image Credits: Xingqin Lin et al.

Rayleigh RWP

The mean transition length and time are as follows:

$$E[L] = \frac{1}{2\sqrt{\lambda}}$$

$$E[T] = \frac{1}{2v\sqrt{\lambda}}$$

Sojourn Time and Handover Rate in cellular networks, Xingqin Lin et al.

 Sojourn Time and Handover Rate in cellular networks, Xingqin Lin et al.
 Node distribution during one movement period

$$f(r,\theta) = \frac{\sqrt{\lambda}}{\pi r} \exp(-\lambda \pi r^2)$$

 Sojourn Time and Handover Rate in cellular networks, Xingqin Lin et al.
 Node distribution during one movement period

$$f(r,\theta) = \frac{\sqrt{\lambda}}{\pi r} \exp(-\lambda \pi r^2)$$

$$S_T = E[T] \int_A f(r,\theta) dA$$

 Sojourn Time and Handover Rate in cellular networks, Xingqin Lin et al.
 Node distribution during one movement period

$$f(r,\theta) = \frac{\sqrt{\lambda}}{\pi r} \exp(-\lambda \pi r^2)$$

$$S_T = E[T] \int_A f(r,\theta) dA$$

 Analysis of Cell Sojourn Time in Heterogeneous Networks With Small Cells, S. Shin et al.
 Expected amount of time a mobile spends in macro-cell-only-area (MoA)

Sojourn Time

If we know the expected number of transitions E[N] in which a node moves out of the region, then sojourn time can be given by

$$S_T = (E[N] - 1)E[T] + E[T_{last}]$$

where T_{last} is the time spent inside the region during the last transition. Since it is difficult to characterize T_{last} ,

$$S_T \approx (E[N] - 1/2)E[T]$$

E[N]

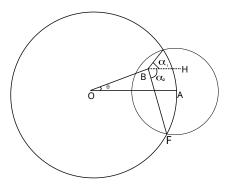
$$E(N) = \sum_{k=1}^{\infty} kPr(r, \theta, k)$$

where

$$Pr(r, \theta, k) = \int_{S-A} \int_{A} \dots$$
$$\int_{A} f_{X_{1}/X_{0}}(x_{1}/x_{0}) \dots f_{X_{k+1}/X_{k}}(x_{k+1}/x_{k}) dA_{1} \dots dA_{k+1}$$

 $f_{X_n/X_{n-1}}(x_n/x_{n-1})$ is the probability density of the destination X_n given that the node's current position is X_{n-1}

Leaving in One Transition



$$\rho(r,\theta) = Pr(-\alpha_2 < \alpha < \alpha_1, r_1 > r_{11}) + Pr(\alpha_1 < \alpha < 2\pi - \alpha_2, r_1 > r_{12})$$

$$= \int_{-\alpha_2}^{\alpha_1} \int_{r_{11}}^{\infty} f_{r_1,\alpha}(r_1,\alpha) dr_1 d\alpha + \int_{\alpha_1}^{2\pi - \alpha_2} \int_{r_{12}}^{\infty} f_{r_1,\alpha}(r_1,\alpha) dr_1 d\alpha$$

This is a general expression that can be used for any mobility model.



In case of RWP, r_1 and α are chosen independently. Therefore, $f_{r_1,\alpha}(r_1,\alpha)=f_{r_1}(r_1)f_{\alpha}(\alpha)$

$$\rho(r,\theta) = \int_{-\alpha_2}^{\alpha_1} f_{\alpha}(\alpha) \int_{r_{11}}^{\infty} f_{r_{1}}(r_{1}) dr_{1} d\alpha + \int_{\alpha_1}^{2\pi - \alpha_2} f_{\alpha}(\alpha) \int_{r_{12}}^{\infty} f_{r_{1}}(r_{1}) dr_{1} d\alpha
= \int_{-\alpha_2}^{\alpha_1} \frac{1}{2\pi} e^{-\lambda \pi r_{11}^2} d\alpha + \int_{\alpha_1}^{2\pi - \alpha_2} \frac{1}{2\pi} e^{-\lambda \pi r_{12}^2} d\alpha$$

Leaving in One Transition

The probability with which a node at (r, θ) moves out of the region of interest during the next transition is

$$\rho(r,\theta) = \int_{-\alpha_2}^{\alpha_1} \frac{1}{2\pi} e^{-\lambda \pi r_{11}^2} d\alpha + \int_{\alpha_1}^{2\pi - \alpha_2} \frac{1}{2\pi} e^{-\lambda \pi r_{12}^2} d\alpha$$

Where

$$r_{11} = -r\cos(\theta - \alpha) + \sqrt{R_0^2 - r^2\sin^2(\theta - \alpha)}$$

$$r_{12} = R_0 \cos \alpha - r \cos(\theta - \alpha) + \sqrt{[R_0 \cos \alpha - r \cos(\theta - \alpha)]^2 - r^2 \sin^2 \theta - [R_0 - r \cos \theta]^2 + R_2^2}$$

Leaving in One Transition

$$\beta_2 = \cos^{-1}\left(1 - \frac{R_2^2}{2R_0^2}\right) \quad \beta_1 = \beta_2 - \theta$$

$$\alpha_1 = \theta + \tan^{-1} \left(\frac{R_0 \sin \beta_1}{R_0 \cos \beta_1 - r} \right)$$

$$\alpha_2 = \tan^{-1} \left(\frac{r \sin \theta + R_0 \sin \beta_2}{R_0 \cos \beta_2 - r \cos \theta} \right)$$

Simulations

For all simulations, a 1000×1000 square is used to represent the whole 2-D plane. $R_0=150, R_2=0.4R_0$

To draw a length from Rayleigh distribution, the following method is used:

- 1. Draw a number N from Poisson distribution with density λA where A is the area of the square.
- 2. Distribute these N points uniformly on the square
- 3. Of these N points, choose the point that is closest to the point under consideration.

This leads to Rayleigh distribution of transition length and can be proved using null probability of a Poisson Point Process.

Simulations

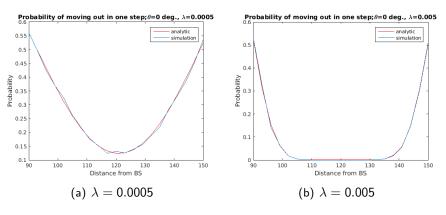
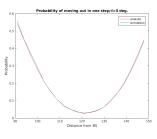
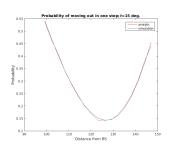


Figure: $\rho(r,0)$ vs. λ

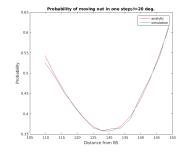
Simulations





(a)
$$\theta = 5^{\circ}$$

(b)
$$\theta=15^\circ$$





E[N]

Let us make a gross approximation and use the same probability of leaving for all waypoints in the path. Then the average number of steps a node starting at (r, θ) takes to leave the region is given by

$$E[N] = \sum_{k=1}^{\infty} k(1 - \rho(r, \theta))^{k-1} \rho(r, \theta)$$
$$= \frac{1}{\rho(r, \theta)}$$

E[N]

The plots are for points along the radius at angles $\theta=10^\circ$ and $\theta=15^\circ$. We can see that the analytical formula agrees better for narrower regions.

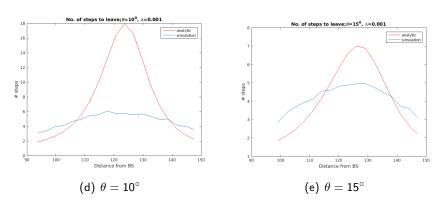


Figure: Expected number of transitions

Ongoing Work Markov Chain Process

The transitions in this mobility model have Markovian property

$$f_{X_n/X_{n-1},X_{n-2},...,X_0}(x_n/x_{n-1},x_{n-2},...,x_0) = f_{X_n/X_{n-1}}(x_n/x_{n-1})$$

Where X_{n-1} is the current waypoint and X_n is the next waypoint.

Ongoing Work

Discretizing the State Space

- ▶ The state space of the Markov chain is continuous
- ▶ Discretise the space into n+1 states n states lie inside the region of interest and the n + 1th state represents space outside the region

Ongoing Work

Discretizing the State Space

- ▶ The state space of the Markov chain is continuous
- ▶ Discretise the space into n+1 states n states lie inside the region of interest and the n + 1th state represents space outside the region
- Model the motion as an Absorbing Markov Chain The transition probabilities among first n states depend on the distances between the states and the transition probabilities from these n states to the absorbing state is $\rho(r_i, \theta_i)$

The transition matrix is

$$P = \left(\begin{array}{cc} Q & R \\ \mathbf{0} & 1 \end{array}\right)$$

 $Q_{n\times n}$ is the transition matrix of n non-absorbing states $R_{n\times 1}$ contains the probabilities of moving out in one step from each of those n states.

Expected number of transitions $\mathbf{t}_{n\times 1}$

$$\mathbf{t} = F\mathbf{1}$$

and the variance vector V is given by

$$V = (2F - I)\mathbf{t} - \mathbf{t}_{sq}$$

where
$$F = \sum_{k=0}^{\infty} Q^k = (I - Q)^{-1}$$

References

- X. Lin, R. K. Ganti, P. J. Fleming, and J. G. Andrews, Towards understanding the fundamentals of mobility in cellular networks, IEEE Transactions on Wireless Communications, vol. 12, 2013.
- S. Shin, U. Lee, F. Dressler and H. Yoon, "Analysis of Cell Sojourn Time in Heterogeneous Networks With Small Cells," IEEE Communications Letters. vol. 20, 2016
- H. Elkotby and M. Vu, Interference and throughput analysis of uplink user-assisted relaying in cellular networks, PIMRC, 2014.
- R. K. Ganti, Stochastic geometry and wireless networks. SPCOM, July 2012.
- Wikipedia, Absorbing Markov Chain wikipedia, the free encyclopedia,.