

Time Reversal of Some Stationary Jump-Diffusion Processes from Population Genetics

Martin Hutzenthaler, Jesse E. Taylor

(Submitted on 15 Nov 2010)

We describe the processes obtained by time reversal of a class of stationary jump-diffusion processes that model the dynamics of genetic variation in populations subject to repeated bottlenecks. Assuming that only one lineage survives each bottleneck, the forward process is a diffusion on $[0,1]$ that jumps to the boundary before diffusing back into the interior. We show that the behavior of the time-reversed process depends on whether the boundaries are accessible to the diffusive motion of the forward process. If a boundary point is inaccessible to the forward diffusion, then time reversal leads to a jump-diffusion that jumps immediately into the interior whenever it arrives at that point. If, instead, a boundary point is accessible, then the jumps off of that point are governed by a weighted local time of the time-reversed process.

Comments: 23 pages, 1 figure, to appear in Advances in Applied Probability Vol 42, No 4, p 1-25

Subjects: **Probability (math.PR)**

MSC classes: 60J60, 60J55, 92D10

Cite as: [arXiv:1011.3379v1](#) [math.PR]

Submission history

From: Martin Hutzenthaler [[view email](#)]

[v1] Mon, 15 Nov 2010 13:42:43 GMT (49kb,D)

[Which authors of this paper are endorsers?](#)

Download:

- [PDF](#)
- [Other formats](#)

Current browse context:

math.PR

[< prev](#) | [next >](#)

[new](#) | [recent](#) | [1011](#)

Change to browse by:

[math](#)

References & Citations

- [NASA ADS](#)

Bookmark([what is this?](#))

