

## Supplemental Information for:

### andscape attributes governing local transmission of an endemic zoonotic virus: rabies in domestic dogs.

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#### METHODS S1

##### 454 pyrosequencing

Whole genome sequences for five samples (see Table S1) were generated via 454 pyrosequencing protocols. DNA for three samples was prepared according to the depletion method detailed in the main methods and sent to the Animal & Plant Health Agency (APHA) sequencing facility for the preparation of sequencing libraries and 454 pyrosequencing. The other two samples were prepared according to a PCR amplicon approach as follows. TRIzol-extracted viral RNA was reverse transcribed to cDNA using the primer RABV\_Tzdg.p1f (5' ACGCTTAACAACAAAATCAGAG 3') at a concentration of 2pmol/ $\mu$ l and Superscript III reverse transcriptase (Invitrogen) in a total volume of 20 $\mu$ l, as per manufacturer's instructions. An in house set of 26 short, tagged, overlapping primer pairs spanning the entire RABV genome was used to obtain PCR products with 1 $\mu$ l of cDNA and

KOD hot start DNA polymerase kit as per the manufacturer's protocol (Novagen). The following cycling parameters were used: 1 hold at 95 °C for 2 mins, 35 cycles at 95 °C for 20 s, 50-60°C (dependent on the optimized temperature for each primer pair) for 20 s, 70 °C for 20 s and a final hold at 70 °C for 10 min. Products were pooled for each sample with libraries and sequencing completed by the APHA sequencing facility.

## **Construction of resistance surfaces**

Landscape attributes were characterized as surface models by assigning cell values to represent the assumed facilitating or impeding impact of a predictor on RABV diffusion. Resistance surfaces for each attribute were formatted as follows:

### i) Dog density

Dog density estimates were taken from a household census conducted in the Serengeti District in 2014-2015, which provided the GPS location and dog count for each household. The density point pattern was smoothed across the default raster grid using an isotropic Gaussian smoothing kernel with  $\sigma=500$  using the R package *spatstat* (1) with cell values expressing the estimated intensity values. As dog density is assumed to facilitate RABV diffusion, the reciprocal of these values was used as a resistance value in each cell. Serengeti National Park (SNP) areas were assigned a low resistance value, equivalent to 1 dog per  $\text{km}^2$  to reflect the low dog density in this area. Cells outside the district (accounting for ~12% of all cells) were assigned random values from the Serengeti density data.

### ii) Dog presence

Dog census data as described above was used to assign a presence or absence value to 100m grid cells based on the exact locations of dogs. Dog presence was assigned a low resistance value of 0.1 to represent conductive "corridors" of movement through normal landscape, whereas absence was the null IBD value of 1 to represent higher resistance to dog movement outside connected corridors.

### iii) Elevation and slope

A digital elevation model covering the landscape was converted to raster format and elevation values used directly as resistance values. Slope values were calculated from the digital elevation model using the SDMTools package (2) in R. All slope values were increased by a value of 1 to ensure comparison to a null IBD surface was possible. The resolution of the digital elevation model was not consistent across the extent of the landscape as some areas have been mapped in more detail, which resulted in some finer grained areas in raster grids.

#### iv) Human to dog ratio (HDR)

Household census data was used to estimate the human to dog ratio per village, which was assigned directly as a resistance value in each cell.

#### v) Major roads and rivers

Shapefiles of major roads and rivers in the study area were converted to raster grids (one for roads, one for rivers) with the defined spatial extent and resolution. Cells containing a road feature were assigned a low resistance value of 0.001 to represent increased diffusion along roads relative to uniform landscape, all other cells were assigned a value of 1. River cells were assigned high resistance values of 1000 to reflect their influence as a barrier to diffusion.

#### vi) Average vaccination coverage

Mass dog vaccination campaigns have been undertaken in the Serengeti District since 2002, with varying annual coverage across villages. We used an average annual % vaccination coverage per village across the 11-year period from 2002 up to 2013 and assigned values to grid cells as resistance values [range: 6.43-100]. Rabies appears to have been locally eliminated in the SNP and there is a requirement for all dogs to be vaccinated within the park boundaries, therefore SNP cells were assigned the highest resistance value of 100. Cells outside the district and SNP were assigned the minimum observed average vaccination coverage of 6.43% as there is no formal vaccination initiative in these areas and therefore coverage is assumed to be low.

#### vi) Susceptible dog density

The dog density estimates were depleted according to the averaged vaccination coverage described above to produce a resistance surface representative of the susceptible host population density.

## vii) Number of vaccination campaigns

The number of vaccination campaigns conducted across the sampled time period 2004-2013 was summed for each village and assigned to each grid cell. Note some villages experienced return visits when turnout was less than expected, these were counted as separate campaigns if a coverage of at least 10% was achieved.

## **Determining an optimum location state partitioning scheme**

Rabies cases projected in MDS attribute-defined space were partitioned according to varying levels of spatial aggregation using a *k*-means algorithm. The following methods were used to obtain an optimum range of *k*-values to determine a partitioning scheme:

- i) Elbow method: the point of maximum curvature in a plot of number of clusters versus within-group sum of squares;
- ii) Partitioning around medoids: using the optimum average silhouette to estimate the number of clusters, R package: *fpc* (3)
- iii) Model-based clustering: chooses the optimal model and number of clusters according to Bayesian Information Criterion for expectation-maximisation, R package: *mclust* (4)
- iv) Affinity propagation: a clustering algorithm that takes a pairwise similarity matrix and simultaneously considers all data points as potential cluster centres. The algorithm finds an optimum set of clusters that maximises the total similarity between data points and their cluster centres by an iterative process (5), R package: *apcluster* (6);
- v) Gap statistic: a statistical procedure to formalise the "elbow" method by comparing the change in within-cluster dispersion to a reference null distribution (7), R package: *cluster* (8);
- vi) R package *NbClust*: provides 30 indices to determine the number of clusters (9).

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**S1 Table. Model selection results of the continuous phylogeography diffusion model for Tanzania rabies virus dataset.** Shown are marginal likelihood estimates using path sampling and stepping stone sampling in BEAST, with the best model in bold.

Diffusion model	Path Sampling	Stepping Stone
Homogeneous	-26674.93	-26679.87
<b>Gamma</b>	<b>-26661.45</b>	<b>-26666.37</b>
Lognormal	-26683.68	-26688.12

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**S2 Table. Pearson's correlation coefficients between paired resistance distance matrices for landscape attributes used to parameterize GLM diffusion models estimating the rabies virus movement in the Serengeti District.** Pairwise resistance distance matrices were constructed for  $k=5$  to 15 centroids. Correlations greater than 0.9 are highlighted in large bold text.

Pairwise attribute combinations	k5	k6	k7	k8	k9	k10	k11	k12	k13	k14	k15
Campaigns,Dog density	0.55	0.85	0.58	0.45	0.56	0.51	0.70	0.54	0.35	0.30	0.39
Campaigns,Dog presence	0.58	0.65	0.66	0.13	0.30	0.12	0.08	0.57	0.26	0.34	0.41
Campaigns,Elevation	0.53	0.50	0.67	0.21	0.36	-0.21	-0.12	0.58	-0.11	0.44	0.23
Campaigns,HDR	0.48	0.42	-0.05	0.63	0.70	-0.53	-0.05	0.62	-0.32	0.17	0.74
Campaigns,Rivers	-0.29	0.10	0.02	-0.13	0.07	-0.22	-0.27	0.15	0.12	0.30	0.00
Campaigns,Roads	0.87	0.74	0.60	0.57	0.56	0.27	0.17	0.63	0.62	0.28	0.02
Campaigns,Slope	-0.01	0.15	-0.15	-0.28	-0.31	-0.67	-0.62	-0.12	-0.27	-0.10	-0.51
Campaigns,Susceptibles	0.52	0.84	0.66	0.53	0.67	0.46	0.68	0.69	0.48	0.39	0.41
Campaigns,VaccCov	0.57	0.70	0.88	0.72	0.77	-0.51	-0.07	0.72	0.73	0.85	0.47
Dog density,Dog presence	0.86	<b>0.91</b>	0.73	0.82	0.75	-0.17	0.06	0.68	0.66	0.63	0.09
Dog density,Elevation	0.47	0.65	0.35	0.44	0.38	0.43	0.28	0.47	-0.05	0.36	0.37
Dog density,HDR	-0.01	0.14	0.10	0.03	0.17	-0.32	-0.03	0.01	-0.08	-0.04	0.28
Dog density,Rivers	0.14	0.34	-0.22	0.41	0.39	-0.10	-0.18	0.27	0.04	0.21	-0.02
Dog density,Roads	0.22	0.69	0.69	0.26	0.37	0.59	-0.19	0.52	0.66	0.51	0.51
Dog density,Slope	0.78	0.61	0.22	0.54	0.21	-0.67	-0.47	0.05	0.35	0.21	-0.54
Dog density,Susceptibles	<b>0.97</b>	<b>0.99</b>	<b>0.98</b>	<b>0.98</b>	<b>0.98</b>	<b>0.96</b>	<b>0.98</b>	<b>0.96</b>	<b>0.96</b>	<b>0.97</b>	<b>0.98</b>
Dog density,VaccCov	0.30	0.55	0.23	0.31	0.36	-0.22	0.28	0.20	0.09	0.40	0.38
Dog presence,Elevation	0.85	0.86	0.77	0.80	0.83	0.11	0.15	<b>0.90</b>	0.00	0.86	0.14
Dog presence,HDR	0.19	0.20	-0.10	0.03	0.20	0.04	0.11	0.38	-0.17	0.03	0.51
Dog presence,Rivers	0.24	0.59	-0.13	0.75	0.78	0.57	0.51	0.73	0.00	0.08	0.55
Dog presence,Roads	0.34	0.66	0.84	0.26	0.44	-0.06	0.26	0.57	0.54	0.63	0.19
Dog presence,Slope	0.67	0.67	0.29	0.76	0.54	0.38	0.43	0.19	0.61	0.56	0.08
Dog presence,Susceptibles	0.88	<b>0.93</b>	0.74	0.82	0.73	-0.16	0.06	0.70	0.64	0.66	0.12

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Dog presence,VaccCov	0.45	0.53	0.51	0.06	0.11	-0.16	0.12	0.36	0.22	0.33	-0.12
Elevation,HDR	0.42	0.55	-0.18	0.39	0.49	0.32	-0.11	0.60	0.65	0.09	0.20
Elevation,Rivers	0.26	0.78	0.10	0.75	0.79	0.25	0.26	0.71	0.35	0.04	0.20
Elevation,Roads	0.52	0.52	0.46	0.49	0.57	0.41	-0.06	0.54	0.03	0.49	0.45
Elevation,Slope	0.28	0.41	0.15	0.47	0.33	0.10	0.30	0.14	0.02	0.36	-0.04
Elevation,Susceptibles	0.51	0.69	0.42	0.52	0.43	0.46	0.33	0.55	-0.08	0.41	0.43
Elevation,VaccCov	0.49	0.55	0.74	0.09	0.17	0.37	0.31	0.48	-0.15	0.39	0.52
HDR,Rivers	0.42	0.32	-0.03	0.10	0.31	0.42	0.24	0.36	0.14	0.05	0.24
HDR,Roads	0.51	0.21	0.05	0.21	0.23	-0.03	-0.21	0.28	-0.11	-0.13	-0.09
HDR,Slope	-0.49	-0.42	0.08	-0.37	-0.48	0.35	0.06	-0.33	-0.10	0.05	-0.05
HDR,Susceptibles	0.14	0.21	0.10	0.15	0.32	-0.16	-0.02	0.21	-0.12	-0.07	0.27
HDR,VaccCov	<b>0.91</b>	0.75	-0.13	0.73	0.79	0.90	0.15	0.80	-0.30	0.10	0.24
Rivers,Roads	-0.29	0.06	-0.40	0.09	0.22	-0.07	0.23	0.33	-0.08	0.16	0.26
Rivers,Slope	0.16	0.51	0.35	0.52	0.35	0.51	0.60	0.25	-0.04	-0.09	0.50
Rivers,Susceptibles	0.28	0.34	-0.28	0.42	0.39	-0.01	-0.14	0.28	-0.06	0.33	0.04
Rivers,VaccCov	0.43	0.10	-0.06	-0.13	0.02	0.24	-0.03	0.13	0.23	0.44	0.01
Roads,Slope	-0.32	0.06	0.04	-0.13	0.00	-0.57	-0.09	-0.07	-0.03	0.03	-0.19
Roads,Susceptibles	0.14	0.70	0.75	0.33	0.45	0.63	-0.11	0.63	0.72	0.60	0.58
Roads,VaccCov	0.40	0.54	0.43	0.02	0.06	0.10	-0.23	0.19	0.36	0.06	0.15
Slope,Susceptibles	0.75	0.57	0.08	0.51	0.11	-0.67	-0.51	-0.09	0.22	0.15	-0.53
Slope,VaccCov	-0.16	-0.16	-0.40	-0.13	-0.36	0.29	-0.25	-0.34	-0.41	-0.11	-0.24
Susceptibles,VaccCov	0.48	0.64	0.36	0.35	0.44	-0.08	0.32	0.37	0.24	0.49	0.45



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**S3 Table. Epidemiological information and whole genome sequencing (WGS) details for 152 rabies virus samples collected from the Serengeti District, Tanzania, between 2004 and 2013.**

Sample ID	Date of sample collection	Easting	Northing	Host	WGS protocol	Total quality trimmed reads	Viral reads mapped	Proportion viral reads	Average depth	GenBank Accession	Details/Reference
RV2483	28/09/08	701590	9794710	Goat	MiSeq	397221	2045	0.51	22.05	KY210220	This paper
RV2485	28/09/08	701590	9794710	Goat	MiSeq	300966	907	0.30	8.12	KY210221	This paper
RV2489	17/01/09	699900	9794413	Domestic dog	MiSeq	2665913	3359	0.13	16.43	KY210222	This paper
RV2490	07/08/08	702122	9796402	Domestic dog	MiSeq	1013605	8526	0.84	79.09	KR534217	Original GenBank submission (Brunker <i>et al</i> , 2015) updated to WGS for this paper
RV2491	02/08/08	696748	9794246	Domestic dog	MiSeq	1421454	10733	0.76	101.98	KR534218	Original GenBank submission (Brunker

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											<i>et al</i> ,2015) updated to WGS for this paper
RV24 92	03/11/07	685 837	9799 418	Dom estic dog	MiSeq	1642925	12895	0.78	100.93	KR53 4219	Original GenBank submissio n (Brunker <i>et al</i> ,2015) updated to WGS for this paper
RV24 93	31/07/08	700 689	9793 979	Dom estic dog	MiSeq	1304543	4780	0.37	44.57	KR53 4220	Original GenBank submissio n (Brunker <i>et al</i> ,2015) updated to WGS for this paper
RV24 95	12/07/08	696 385	9802 796	Dom estic dog	MiSeq	1946750	14768	0.76	129.48	KR90 6734	Brunker <i>et al</i> ,2015

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RV24 97	22/06/07	678 577	9807 715	Dom estic dog	MiSeq/NextSeq	15591341	6031	0.04	63.98	KY21 0223	This paper
RV25 00	23/11/08	698 019	9804 712	Dom estic dog	MiSeq	557886	1536	0.28	15.86	KR90 6737	Brunker <i>et al</i> ,2015
RV25 01	14/02/04	679 372	9810 508	Dom estic dog	MiSeq	1030023	11539	1.12	108.23	KR90 6738	Brunker <i>et al</i> ,2015
RV25 03	15/02/09	686 729	9761 151	Wild cat	MiSeq	1325075	2774	0.21	28.64	KR90 6740	Brunker <i>et al</i> ,2015
RV27 67	12/09/09	665 178	9777 064	Dom estic dog	MiSeq	1039525	23560	2.27	249.79	KR90 6742	Brunker <i>et al</i> ,2015
RV27 86	20/06/10	670 876	9811 501	Dom estic dog	MiSeq/NextSeq	15706254	10705	0.07	109.00	KY21 0224	This paper
RV27 88	05/08/10	668 396	9794 012	Dom estic dog	MiSeq	1952400	16416	0.84	181.53	KR53 4228	Original GenBank submissio n (Brunker <i>et al</i> ,2015) updated to WGS for this paper
RV27 89	22/10/10	686 741	9795 339	Dom estic	MiSeq	825552	1753	0.21	16.94	KR53 4229	Original GenBank

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				dog							submissio n (Brunker <i>et al</i> ,2015) updated to WGS for this paper
RV27 90	01/11/10	678 362	9807 838	Dom estic dog	MiSeq	190118	2054	1.08	20.87	KR53 4230	Original GenBank submissio n (Brunker <i>et al</i> ,2015) updated to WGS for this paper
RV27 91	08/11/10	681 329	9802 616	Dom estic dog	MiSeq/NextSeq	15068608	20740	0.14	237.54	KR53 4231	Original GenBank submissio n (Brunker <i>et al</i> ,2015) updated to WGS for this paper

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RV27 92	07/01/11	670 624	9824 613	Jackal	MiSeq/NextSeq	13995380	19366	0.14	215.26	KR53 4232	Original GenBank submissio n (Brunker <i>et al</i> ,2015) updated to WGS for this paper
RV27 93	02/01/11	657 176	9822 908	Cow	MiSeq	149085	3900	2.62	38.81	KR90 6755	Brunker <i>et al</i> ,2015
RV27 94	12/01/11	679 391	9807 221	Domestic dog	MiSeq	514797	2064	0.40	19.28	KR53 4233	Original GenBank submissio n (Brunker <i>et al</i> ,2015) updated to WGS for this paper
RV27 95	16/01/11	682 879	9801 080	Cow	MiSeq	369231	12824	3.47	134.33	KR53 4234	Original GenBank submissio n (Brunker <i>et al</i>

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											,2015) updated to WGS for this paper
RV27 96	19/01/11	671 651	9801 501	Cow	MiSeq	1033943	56028	5.42	508.13	KR53 4235	Original GenBank submissio n (Brunker <i>et al</i> ,2015) updated to WGS for this paper
RV27 97	30/01/11	663 782	9800 181	Cow	MiSeq	1278027	25740	2.01	229.21	KR53 4236	Original GenBank submissio n (Brunker <i>et al</i> ,2015) updated to WGS for this paper
RV27 98	31/01/11	663 782	9800 181	Cow	MiSeq	1287235	69613	5.41	604.45	KR53 4237	Original GenBank submissio n

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											(Brunker <i>et al</i> , 2015) updated to WGS for this paper
RV2799	29/01/11	680129	9805797	Domestic dog	MiSeq	1347238	4352	0.32	41.33	KR906756	Brunker <i>et al</i> , 2015
RV2800	17/02/11	666237	9778079	Domestic dog	MiSeq	142582	2917	2.05	33.48	KR534238	Original GenBank submission (Brunker <i>et al</i> , 2015) updated to WGS for this paper
RV2858	21/02/11	646139	9824316	Goat	MiSeq	1503125	120336	8.01	935.41	KY210225	This paper
RV2859	01/03/11	650148	9819597	Domestic dog	MiSeq	283796	8437	2.97	88.54	KY210226	This paper
RV2861	11/05/11	691423	9791388	Domestic dog	MiSeq	1359627	164027	12.06	1259.37	KR906767	Brunker <i>et al</i> , 2015
RV2862	11/05/11	674259	9805687	Domestic	MiSeq	1177250	2121	0.18	21.18	KR906768	Brunker <i>et al</i>

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				dog							,2015
RV28 63	19/04/11	672 066	9805 512	Dom estic dog	MiSeq	448320	24262	5.41	194.02	KY21 0227	This paper
RV28 66	15/04/11	678 461	9808 213	Goat	MiSeq/NextSeq	17707815	1007	0.01	9.98	KY21 0228	This paper
RV28 67	22/05/11	677 206	9802 038	Dom estic dog	454 pyrosequencing: depletion	2965005	2967	0.10	18.65	KY21 0229	This paper
RV28 68	06/06/11	678 588	9807 873	Cow	MiSeq	703095	27276	3.88	272.91	KY21 0230	This paper
RV28 70	01/06/11	697 216	9792 726	Dom estic dog	MiSeq	529286	19381	3.66	154.52	KY21 0231	This paper
RV28 71	17/06/11	701 315	9794 641	Dom estic dog	454 pyrosequencing: amplicon	46796	8850	18.91	25.34	KR90 6769	Brunker <i>et al</i> ,2015
RV28 73	20/06/11	687 042	9796 792	Dom estic dog	MiSeq	2235772	35378	1.58	359.16	KY21 0232	This paper
RV28 75	29/06/11	680 196	9811 835	Dom estic dog	MiSeq	407187	9927	2.44	99.27	KR90 6770	
RV28 77	30/06/11	679 837	9808 655	Goat	MiSeq	4169008	168447	4.04	1494.91	KY21 0233	This paper
RV28 78	03/07/11	689 579	9796 947	Dom estic dog	MiSeq	146105	1933	1.32	21.85	KY21 0234	This paper
RV28 79	10/06/11	698 379	9806 085	Dom estic dog	MiSeq	696671	6806	0.98	68.17	KY21 0235	This paper



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RV28 80	02/07/11	678 588	9807 873	Cow	MiSeq	795216	8731	1.10	92.11	KY21 0236	This paper
RV28 81	07/07/11	698 357	9806 963	Cow	MiSeq	1101481	12573	1.14	117.68	KY21 0237	This paper
RV28 82	23/07/11	676 741	9807 751	Dom estic dog	454 pyrosequencing: depletion	2896363	35151	1.21	88.58	KY21 0238	This paper
RV28 83	23/07/11	686 525	9797 308	Dom estic dog	MiSeq/NextSeq	20693938	21557	0.10	221.41	KY21 0239	This paper
RV28 84	21/07/11	675 366	9806 739	Cow	MiSeq	1107607	9926	0.90	102.65	KY21 0240	This paper
RV28 85	24/07/11	694 855	9798 730	Goat	MiSeq	969814	3303	0.34	33.06	KY21 0241	This paper
RV28 86	21/07/11	701 234	9794 942	Goat	454 pyrosequencing: amplicon	48056	16226	33.76	64.02	KY21 0242	This paper
RV28 87	25/07/11	695 864	9798 032	Dom estic dog	MiSeq	1044324	2444	0.23	24.46	KY21 0243	This paper
RV28 88	29/07/11	700 262	9793 490	Dom estic dog	MiSeq	1517717	19413	1.28	189.29	KY21 0244	This paper
RV28 89	03/08/11	693 407	9791 698	Dom estic dog	MiSeq	1395952	5503	0.39	50.33	KR53 4244	Original GenBank submissio n (Brunker <i>et al</i> ,2015) updated

# MOLECULAR ECOLOGY

											to WGS for this paper
RV28 90	03/08/11	651 666	9819 798	Cow	MiSeq	1725539	16406	0.95	155.06	KR53 4245	Original GenBank submissio n (Brunker <i>et al</i> ,2015) updated to WGS for this paper
RV28 91	10/08/11	680 536	9813 026	Civet	MiSeq	1290933	54121	4.19	414.53	KR53 4246	Original GenBank submissio n (Brunker <i>et al</i> ,2015) updated to WGS for this paper
RV28 92	12/08/11	676 741	9807 751	Dom estic dog	454 pyrosequencing: depletion	2796394	50188	1.79	91.09	KR53 4247	Original GenBank submissio n (Brunker <i>et al</i>

# MOLECULAR ECOLOGY

											,2015) updated to WGS for this paper
RV28 93	01/08/11	651 666	9819 798	Cow	MiSeq	1637158	31071	1.90	278.08	KR53 4248	Original GenBank submissio n (Brunker <i>et al</i> ,2015) updated to WGS for this paper
RV28 94	15/08/11	653 024	9822 353	Dom estic dog	MiSeq	2125096	188049	8.85	1469.55	KR90 6771	Brunker <i>et al</i> ,2015
RV28 95	18/08/11	678 334	9797 356	Dom estic dog	MiSeq	2273420	19720	0.87	210.02	KR53 4249	Original GenBank submissio n (Brunker <i>et al</i> ,2015) updated to WGS for this paper
RV28	19/08/11	681	9798	Dom	MiSeq	1104977	7214	0.65	68.67	KR90	Brunker

# MOLECULAR ECOLOGY

96		869	035	estic dog						6772	<i>et al</i> ,2015
RV28 97	02/09/11	678 907	9808 303	Dom estic dog	MiSeq	984501	9581	0.97	92.68	KR53 4250	Original GenBank submissio n (Brunker <i>et al</i> ,2015) updated to WGS for this paper
RV28 98	18/09/11	658 269	9804 143	Dom estic dog	MiSeq/NextSeq	19197193	20664	0.11	206.05	KR53 4251	Original GenBank submissio n (Brunker <i>et al</i> ,2015) updated to WGS for this paper
RV28 99	23/09/11	658 245	9804 462	Dom estic dog	MiSeq	897751	8818	0.98	85.53	KR53 4252	Original GenBank submissio n (Brunker <i>et al</i> ,2015)

# MOLECULAR ECOLOGY

											updated to WGS for this paper
RV2900	27/09/11	656396	9803751	Domestic dog	MiSeq	1326869	8924	0.67	84.32	KR906773	Brunker <i>et al</i> , 2015
RV2901	22/09/11	653179	9802910	Domestic dog	MiSeq	522728	10710	2.05	106.87	KR906774	Brunker <i>et al</i> , 2015
RV2902	24/09/11	684532	9790009	Domestic dog	MiSeq	1204863	5234	0.43	43.68	KR906775	Brunker <i>et al</i> , 2015
RV2903	05/10/11	657232	9804521	Domestic dog	MiSeq	1219764	6605	0.54	62.45	KR534253	Original GenBank submission (Brunker <i>et al</i> , 2015) updated to WGS for this paper
RV2906	08/09/11	681338	9808012	Cat	MiSeq	1283722	19254	1.50	170.85	KR534254	Original GenBank submission (Brunker <i>et al</i> , 2015)

# MOLECULAR ECOLOGY

											updated to WGS for this paper
RV2907	16/10/11	700560	9803728	Cow	MiSeq	1353429	7867	0.58	71.47	KR906776	Brunker <i>et al</i> , 2015
RV2909	12/11/11	663440	9804477	Cow	MiSeq	893349	20317	2.27	194.08	KR534256	Brunker <i>et al</i> , 2015
RV3047	10/08/12	655560	9809040	Domestic dog	NextSeq	14822146	193963	1.31	1634.14	KY210245	This paper
RV3048	17/09/12	690718	9803251	Sheep	NextSeq	13774437	92882	0.67	865.92	KY210246	This paper
RV3049	04/10/12	689087	9797146	Domestic dog	NextSeq	14925758	68578	0.46	634.13	KY210247	This paper
RV3050	02/09/12	671593	9804068	Domestic dog	NextSeq	12716032	689856	5.43	5022.73	KY210248	This paper
RV3051	12/10/12	684769	9779316	Domestic dog	NextSeq	13274035	33398	0.25	308.65	KY210249	This paper
RV3052	13/11/12	643810	9813739	Domestic dog	NextSeq	10917330	110185	1.01	1077.20	KY210250	This paper
RV3053	10/11/12	646262	9815621	Domestic dog	NextSeq	11787802	61379	0.52	600.13	KY210251	This paper
RV30	08/12/12	655	9810	Dom	NextSeq	13502719	629341	4.66	4329.86	KY21	This paper

# MOLECULAR ECOLOGY

57		999	468	estic dog						0252	
RV30 58	18/01/13	648 713	9819 325	Dom estic dog	NextSeq	11873546	143723	1.21	1031.42	KY21 0253	This paper
RV30 59	11/01/13	654 132	9825 734	Dom estic dog	NextSeq	12420277	786476	6.33	4587.68	KY21 0254	This paper
RV30 60	04/01/13	654 509	9823 226	Dom estic dog	NextSeq	12150398	316716	2.61	2450.02	KY21 0255	This paper
RV30 61	08/01/13	651 604	9821 249	Dom estic dog	NextSeq	12380235	386682	3.12	2883.85	KY21 0256	This paper
RV30 63	08/01/13	691 639	9810 370	Dom estic dog	NextSeq	11500684	37364	0.32	365.20	KY21 0257	This paper
RV30 64	02/01/13	687 567	9796 857	Dom estic dog	NextSeq	11829231	48612	0.41	469.26	KY21 0258	This paper
RV30 66	23/02/13	652 145	9807 506	Cat	NextSeq	10441152	159856	1.53	1487.65	KY21 0259	This paper
RV30 67	25/02/13	653 061	9777 091	Dom estic dog	NextSeq	15872978	3697	0.02	35.46	KY21 0260	This paper
RV30 68	11/04/13	696 403	9791 890	Goat	NextSeq	11386195	304247	2.67	2459.48	KY21 0261	This paper
RV30 70	27/02/13	673 866	9807 701	Goat	NextSeq	10573869	50029	0.47	504.04	KY21 0262	This paper
RV30 71	13/04/13	680 151	9814 868	Dom estic	NextSeq	13361066	1493699	11.18	7981.02	KY21 0263	This paper

# MOLECULAR ECOLOGY

				dog							
RV30 72	10/04/13	672 356	9817 934	Dom estic dog	NextSeq	10909432	1088443	9.98	6940.73	KY21 0264	This paper
RV30 73	18/02/13	697 846	9808 299	Dom estic dog	NextSeq	9431525	84544	0.90	816.23	KY21 0265	This paper
RV30 74	01/02/13	658 498	9825 235	Dom estic dog	NextSeq	13487972	780515	5.79	5369.35	KY21 0266	This paper
RV30 75	31/01/13	658 498	9825 235	Dom estic dog	NextSeq	12309563	85376	0.69	834.07	KY21 0267	This paper
RV30 78	26/05/13	684 594	9779 133	Cow	NextSeq	13267270	324339	2.44	2550.76	KY21 0268	This paper
RV30 79	18/07/13	700 214	9793 780	Dom estic dog	NextSeq	13140710	10369	0.08	103.78	KY21 0269	This paper
RV30 80	17/07/13	700 300	9793 350	Dom estic dog	NextSeq	12398079	71735	0.58	581.75	KY21 0270	This paper
RV30 82	03/09/13	670 878	9795 201	Dom estic dog	NextSeq	13023573	46367	0.36	485.26	KY21 0271	This paper
RV30 84	22/08/13	697 865	9803 607	Dom estic dog	NextSeq	9206525	18253	0.20	179.18	KY21 0272	This paper
RV30 85	06/09/13	683 848	9810 611	Dom estic dog	NextSeq	11789633	64210	0.54	590.80	KY21 0273	This paper



# MOLECULAR ECOLOGY

RV30 86	06/09/13	697 902	9792 902	Cow	NextSeq	13831121	298860	2.16	2439.13	KY21 0274	This paper
RV30 87	06/09/13	699 819	9795 099	Cow	NextSeq	17902151	12527	0.07	126.18	KY21 0275	This paper
RV30 88	25/04/13	691 229	9812 934	Cow	NextSeq	18221434	358845	1.97	2936.54	KY21 0276	This paper
RV30 90	25/11/11	696 825	9790 124	Cow	MiSeq	129980	10825	8.33	104.41	KY21 0277	This paper
RV30 91	05/12/11	696 698	9792 023	Dom estic dog	MiSeq/NextSeq	18376860	50860	0.28	496.86	KR90 6777	Brunker <i>et al</i> ,2015
RV30 92	16/12/11	680 305	9798 935	Dom estic dog	MiSeq	1045220	2104	0.20	18.69	KY21 0278	This paper
RV30 93	22/12/11	681 720	9798 404	Dom estic dog	MiSeq	495576	4371	0.88	42.70	KR90 6778	Brunker <i>et al</i> ,2015
RV30 94	21/12/11	699 127	9791 922	Dom estic dog	MiSeq	1100604	3942	0.36	35.79	KY21 0279	This paper
RV30 96	30/01/12	658 200	9804 251	Dom estic dog	MiSeq	758254	2900	0.38	24.44	KY21 0280	This paper
RV30 97	19/01/12	658 323	9804 533	Dom estic dog	MiSeq	384498	12069	3.14	113.75	KY21 0281	This paper
RV30 98	26/01/12	658 761	9804 127	Dom estic dog	MiSeq	277003	7448	2.69	73.18	KY21 0282	This paper
RV30 99	19/01/12	696 747	9790 182	Cow	MiSeq	1440888	30135	2.09	233.29	KY21 0283	This paper

# MOLECULAR ECOLOGY

RV3100	19/02/12	698140	9804256	Domestic dog	MiSeq	668576	2648	0.40	24.11	KR906779	Brunker <i>et al</i> , 2015
RV3101	06/03/12	695255	9803449	Jackal	MiSeq	416501	10348	2.48	86.78	KY210284	Brunker <i>et al</i> , 2015
RV3102	08/03/12	700597	9803335	Goat	MiSeq	291477	13309	4.57	123.32	KY210285	Brunker <i>et al</i> , 2015
RV3103	10/12/11	653049	9821771	Domestic dog	MiSeq/NextSeq	17549986	11210	0.06	109.93	KY210286	Brunker <i>et al</i> , 2015
RV3104	22/12/11	648913	9823033	sheep	MiSeq	1145840	11000	0.96	95.11	KR906780	Brunker <i>et al</i> , 2015
RV3105	22/03/12	661920	9803801	Domestic dog	MiSeq	1009153	1365	0.14	13.11	KY210287	This paper
RV3107	09/04/12	656704	9803322	Domestic dog	MiSeq	959083	31920	3.33	292.64	KR906781	Brunker <i>et al</i> , 2015
RV3109	12/04/12	656691	9803564	Domestic dog	MiSeq	409667	24684	6.03	259.04	KY210288	This paper
RV3110	30/04/12	686063	9798923	Domestic dog	MiSeq	346269	6180	1.78	65.54	KY210289	This paper
RV3111	29/04/12	650984	9807685	Domestic dog	MiSeq	897329	2674	0.30	26.85	KR906782	
RV31	06/05/12	685	9792	Dom	MiSeq/NextSeq	18115513	68797	0.38	761.18	KY21	This paper

# MOLECULAR ECOLOGY

13		149	574	estic dog						0290	
RV31 14	06/05/12	685 205	9792 549	Jacka l	MiSeq	854689	16344	1.91	159.71	KY21 0291	This paper
RV31 15	11/05/12	680 871	9782 055	Dom estic dog	MiSeq	800250	3034	0.38	29.22	KY21 0292	This paper
RV31 17	12/05/12	669 765	9795 977	Dom estic dog	MiSeq	2331749	6102	0.26	57.60	KR90 6783	Brunker <i>et al</i> ,2015
RV31 19	17/05/12	680 871	9782 055	Dom estic dog	MiSeq/NextSeq	15519701	7634	0.05	81.87	KY21 0293	This paper
RV31 21	28/05/12	655 915	9809 872	Dom estic dog	MiSeq	1321266	14930	1.13	132.03	KY21 0294	This paper
RV31 22	14/05/12	662 827	9818 718	Dom estic dog	MiSeq	1328914	5972	0.45	56.12	KY21 0295	This paper
RV31 23	26/04/12	647 442	9799 177	Dom estic dog	MiSeq	1530487	7221	0.47	65.08	KR90 6792	Brunker <i>et al</i> ,2015
RV31 24	07/05/12	657 028	9804 565	Dom estic dog	MiSeq	903958	9073	1.00	86.74	KY21 0296	This paper
RV31 25	06/06/12	658 003	9809 433	Goat	MiSeq/NextSeq	16181484	10504	0.06	105.89	KR90 6784	Brunker <i>et al</i> ,2015
RV31 27	11/06/12	681 576	9797 817	Dom estic dog	MiSeq	461052	20729	4.50	204.27	KR90 6785	Brunker <i>et al</i> ,2015

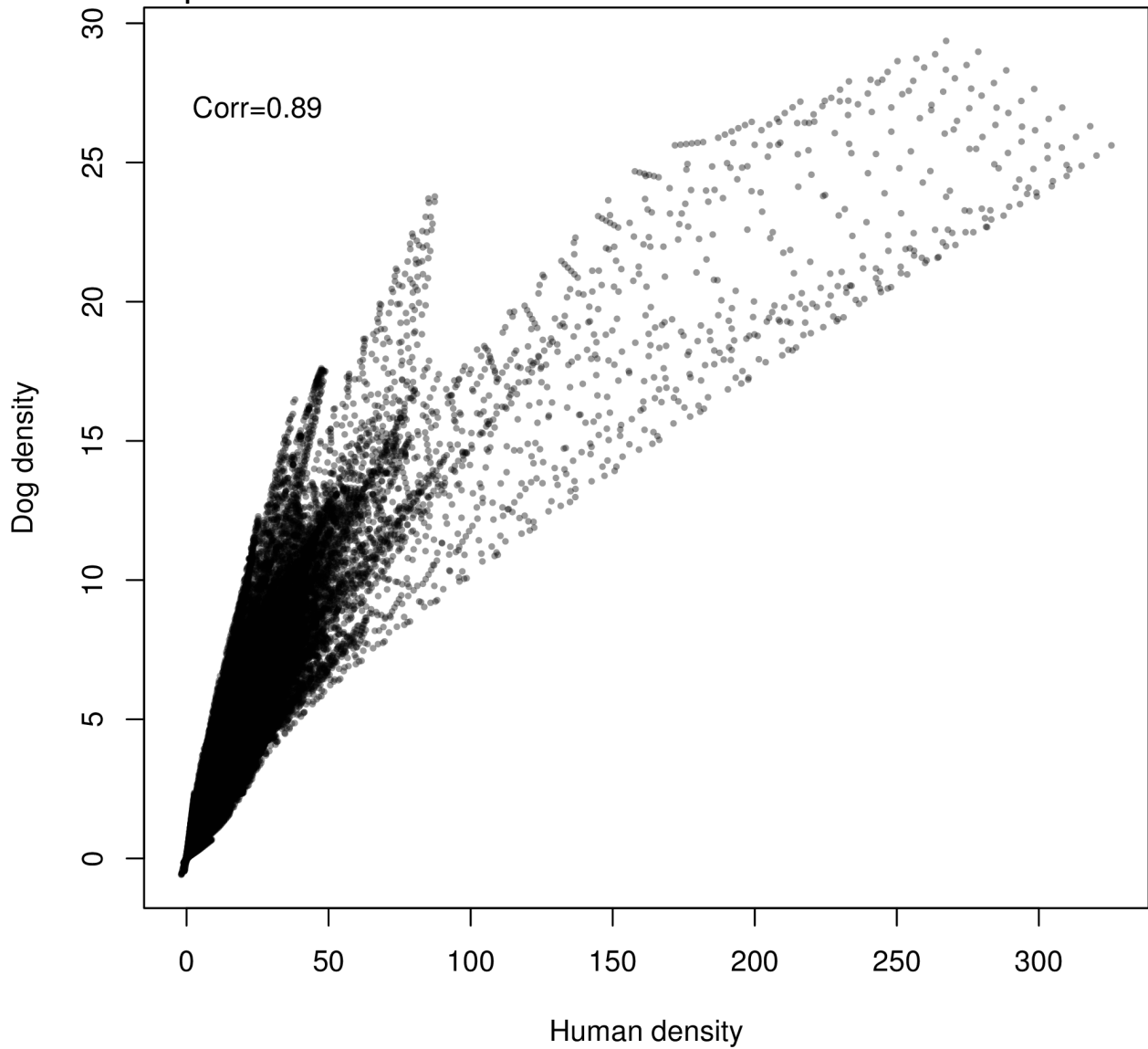
# MOLECULAR ECOLOGY

RV31 28	16/06/12	674 458	9797 076	Donk ey	MiSeq	2044844	9514	0.47	88.78	KR90 6786	Brunker <i>et al</i> ,2015
RV31 29	25/06/12	662 827	9819 718	Shee p	MiSeq	871812	41446	4.75	408.32	KY21 0297	This paper
RV31 30	28/06/12	662 827	9818 718	Goat	MiSeq	155506	10100	6.49	103.35	KY21 0298	This paper
RV31 31	07/07/12	685 498	9797 037	Dom estic dog	MiSeq	1518274	12181	0.80	113.24	KR90 6787	Brunker <i>et al</i> ,2015
RV31 32	02/04/12	656 028	9801 566	Dom estic dog	MiSeq	489623	1478	0.30	13.93	KR90 6788	Brunker <i>et al</i> ,2015
RV31 33	04/07/12	700 856	9800 221	Dom estic dog	MiSeq/NextSeq	14589139	11715	0.08	118.01	KR90 6789	Brunker <i>et al</i> ,2015
RV31 34	19/07/12	689 067	9791 597	Dom estic dog	MiSeq	1280905	3509	0.27	34.25	KY21 0299	This paper
RV31 35	19/07/12	695 591	9788 282	Dom estic dog	MiSeq	1327972	10241	0.77	95.70	KY21 0300	This paper
RV31 36	16/07/12	678 119	9799 572	Cow	MiSeq	993550	15864	1.60	153.90	KY21 0301	This paper
RV31 37	21/07/12	682 623	9783 025	Goat	MiSeq	673235	14221	2.11	140.50	KY21 0302	This paper
RV31 38	16/07/12	677 206	9805 047	Dom estic dog	MiSeq	1634873	13081	0.80	126.10	KY21 0303	This paper
RV31 39	07/07/12	685 498	9797 937	Dom estic	MiSeq	1500822	7764	0.52	74.40	KY21 0304	This paper

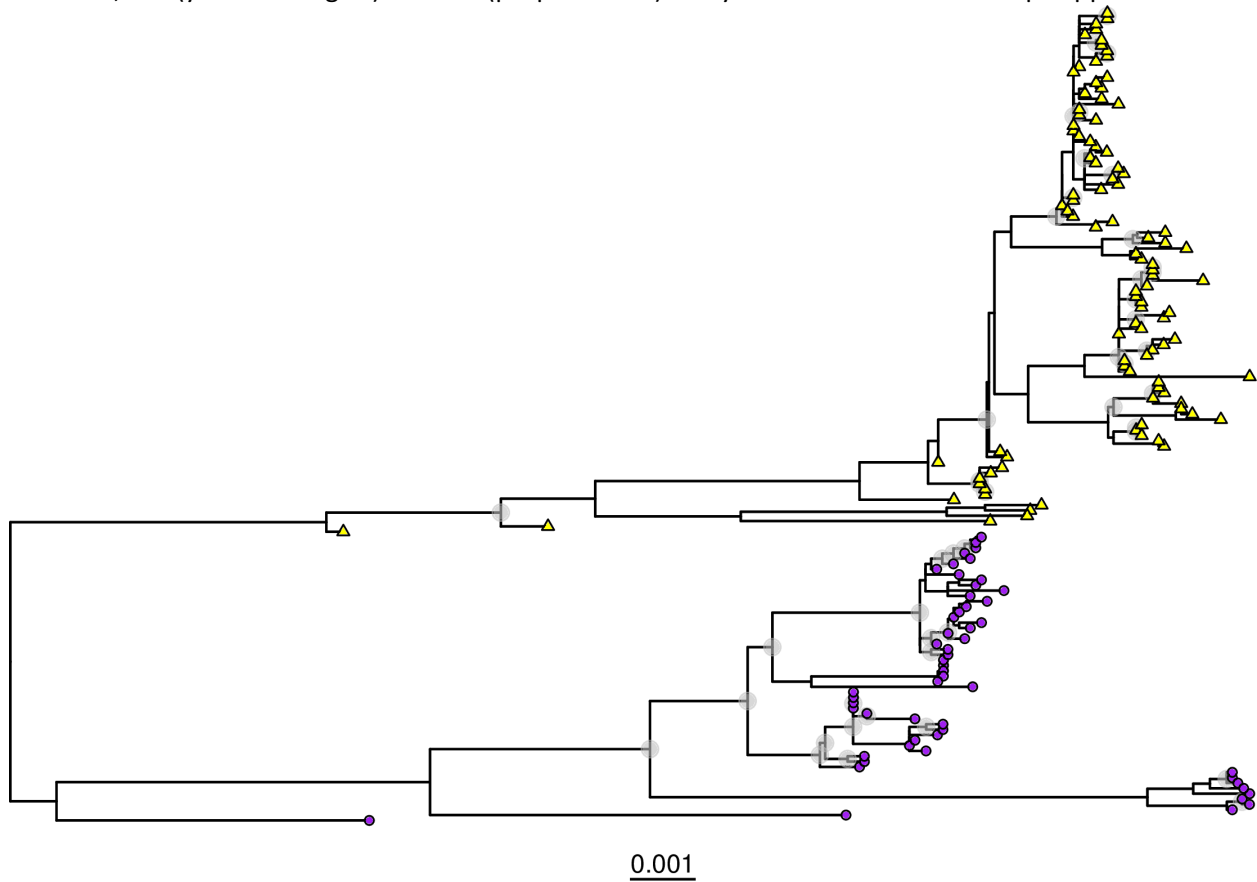
# MOLECULAR ECOLOGY

				dog							
RV31 40	27/07/12	656 945	9804 374	Dom estic dog	MiSeq	1907863	99500	5.22	737.59	KR90 6790	Brunker <i>et al</i> ,2015
RV31 45	08/08/12	658 701	9825 071	Hyen a	MiSeq/NextSeq	15216797	10232	0.07	110.25	KY21 0305	This paper
RV31 46	26/05/12	697 865	9803 637	Civet	MiSeq/NextSeq	16819682	13702	0.08	135.45	KY21 0306	This paper
RV31 49	23/01/12	657 742	9809 576	Dom estic dog	MiSeq	365335	802	0.22	6.23	KR90 6791	This paper
RV31 50	24/03/12	682 197	9791 269	Dom estic dog	MiSeq	1194163	43373	3.63	407.75	KY21 0307	This paper
RV31 51	07/01/12	648 913	9823 033	Cow	MiSeq	105666	6227	5.89	60.67	KY21 0308	This paper
RV31 52	31/05/12	695 002	9803 207	NA	MiSeq/NextSeq	14910632	4003	0.03	42.77	KY21 0309	This paper
RV31 53	23/01/12	694 625	9806 377	Goat	NextSeq	12054558	62937	0.52	637.73	KY21 0310	This paper
RV31 54	29/04/12	658 670	9802 617	Dom estic dog	NextSeq	14618752	401835	2.75	3228.77	KY21 0311	This paper

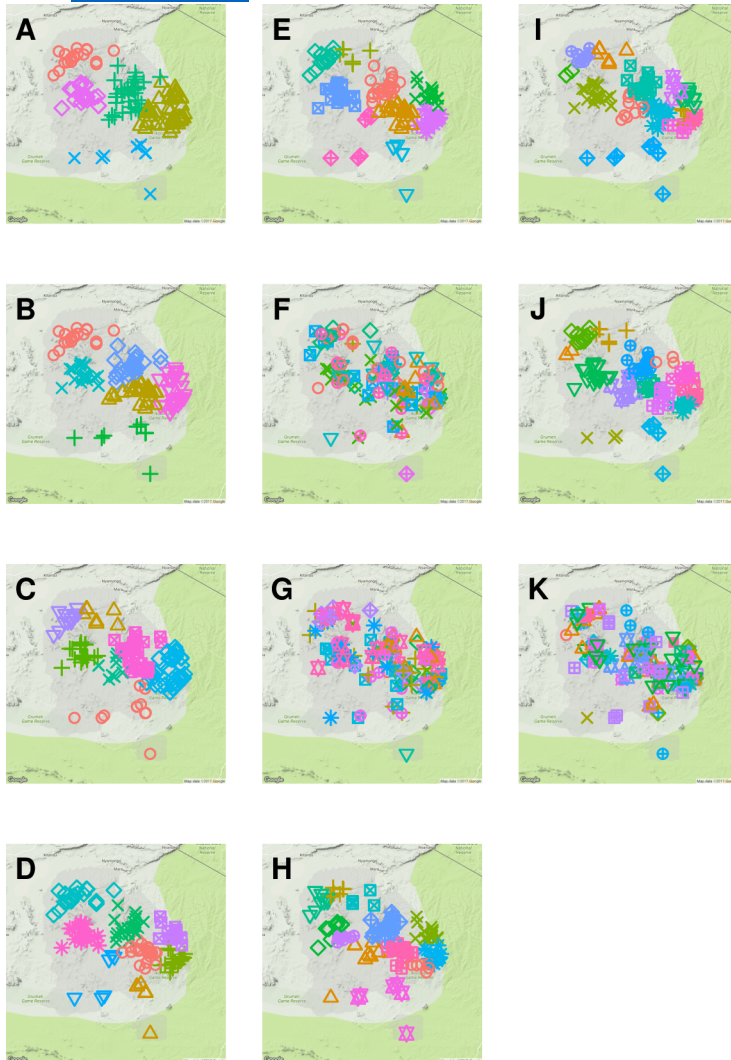
**S1 Fig.** Pearson's correlation between human and dog population density in the Serengeti District based on a complete census conducted in the district.



**S2 Fig. Maximum likelihood phylogeny of 152 rabies cases collected from the Serengeti District, Tanzania 2004-2013.** The phylogeny is scaled in substitutions per site and two major lineages are coloured, Tz1 (yellow triangles) and Tz3 (purple circles). Grey circles indicate bootstrap support  $\geq 80\%$ .



**S3 Fig. Spatial clusters of 152 rabies cases in the Serengeti District according to k-means partitioning using a Euclidean distance matrix.** (A-K) Each cluster is designated by a unique colour/symbol combination for a number of clusters ( $k$ ) ranging from  $k=5$  to  $k=15$ . Map created using R packages ggmap (Kahle & Wickham, 2013) and maptools (Lewin-Koh et al., 2012) with administrative boundaries from [www.nbs.go.tz](http://www.nbs.go.tz).





**S4 Fig. Centroids of spatial clusters assigned according to k-means partitioning using a Euclidean distance matrix for 152 rabies cases in the Serengeti District. (A-K) Centroid positions for a number of clusters ( $k$ ) ranging from  $k=5$  to  $k=15$ . Map created using R package maptools and administrative boundaries from [www.nbs.go.tz](http://www.nbs.go.tz).**

