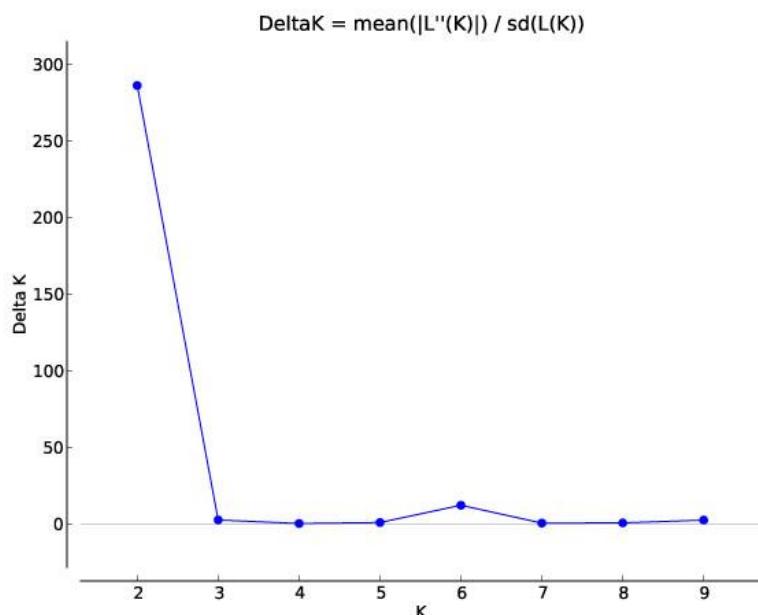


APPENDIX 4 . Results of STRUCTURE analysis for 126 samples belonging to the Guinean group

APPENDIX 4-1. Evanno table output and population assignment for K=1 to K=10

K	Reps	Mean LnP(K)	Stdev LnP(K)	Ln'(K)	Ln''(K)	Delta K
1	20	-5312.27	0.689088	NA	NA	NA
2	20	-4994.44	0.835023	317.83	239.025	286.249652
3	20	-4915.635	14.305695	78.805	37.36	2.611547
4	20	-4799.47	26.294048	116.165	8.71	0.331254
5	20	-4674.595	44.556487	124.875	42.48	0.953397
6	20	-4592.2	5.059228	82.395	61.945	12.243963
7	20	-4571.75	22.797795	20.45	13.655	0.598961
8	20	-4537.645	14.925763	34.105	10.81	0.724251
9	20	-4492.73	10.70062	44.915	27.305	2.551721
10	20	-4475.12	19.313388	17.61	NA	NA



This document was produced by the Structure Harvester, <http://taylor0.biology.ucla.edu/structureHarvester>

CITATION: Earl, Dent A. and vonHoldt, Bridgett M. (2012). STRUCTURE HARVESTER: a website and program for visualizing STRUCTURE output and implementing the Evanno method. Conservation Genetics Resources vol. 4 (2) pp. 359-361 doi: 10.1007/s12686-011-9548-7

Core version: vA.2 July 2014 ; Plot version: vA.1 November 2012 ; Web version: v0.6.94 July 2014;

File generated at 2018-Sep-3 10:48:11

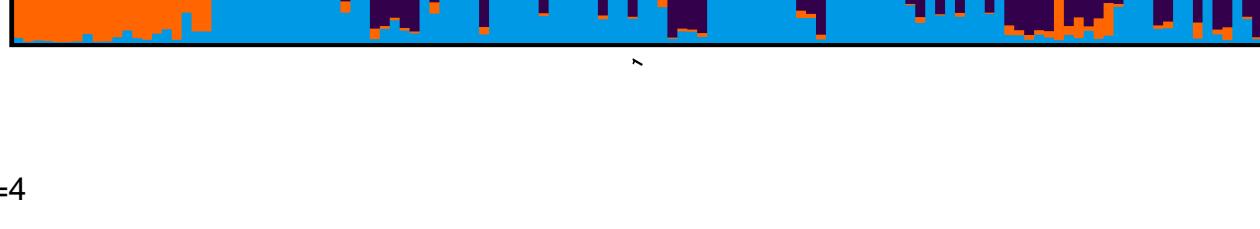
CLUMPAK main pipeline - Job 1533283784 summary

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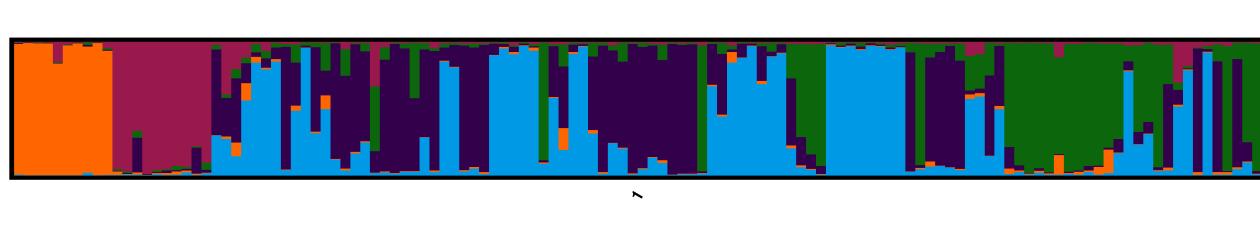
K=1



K=2



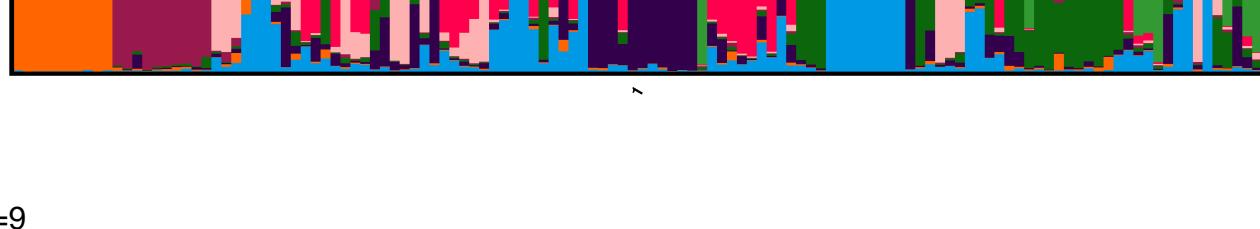
K=3



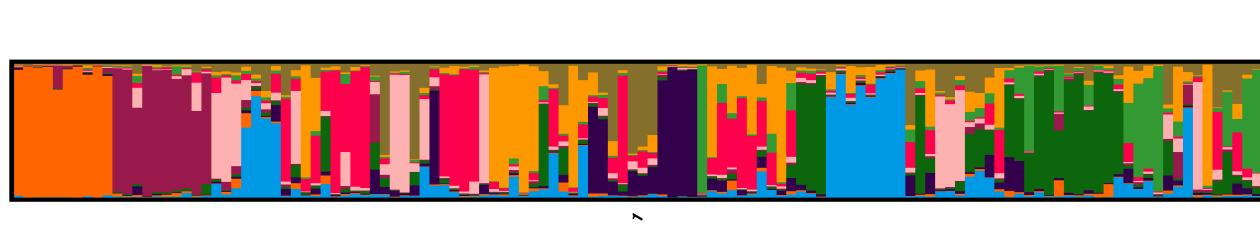
K=4



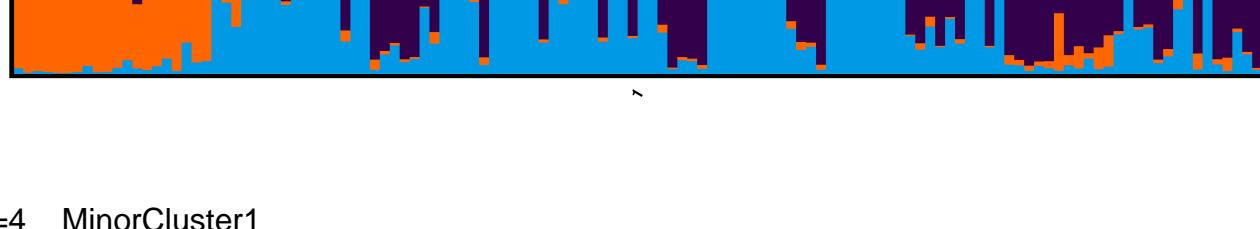
K=5



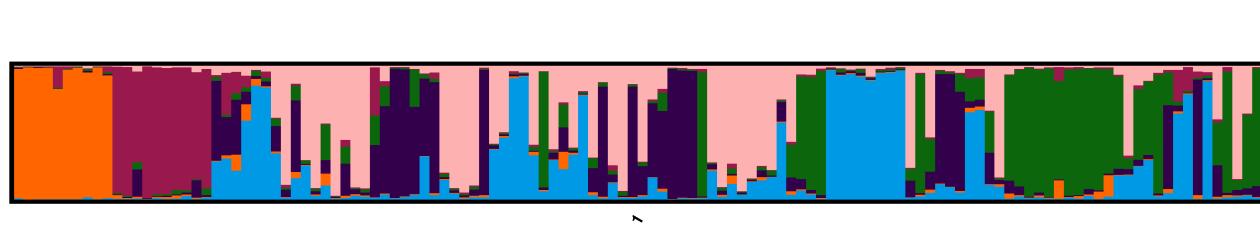
K=6



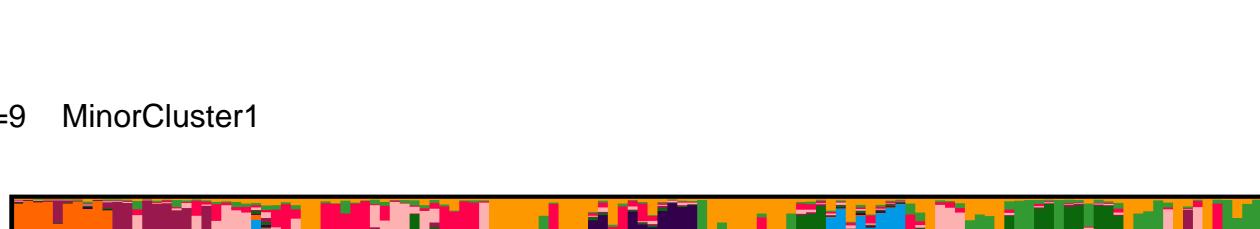
K=7



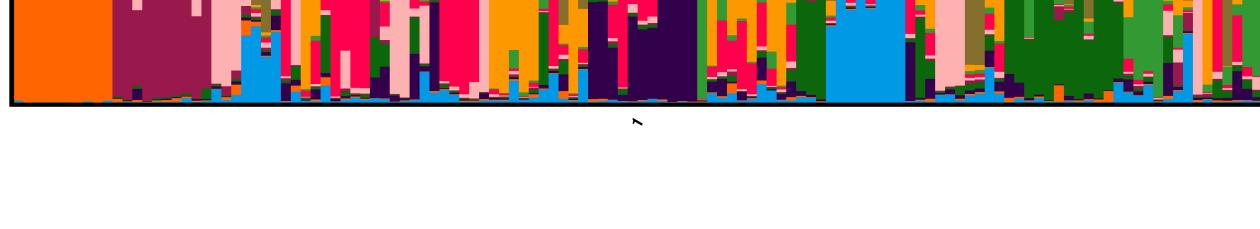
K=8



K=9



K=10



Minor modes for the uploaded data:

K=3 MinorCluster1



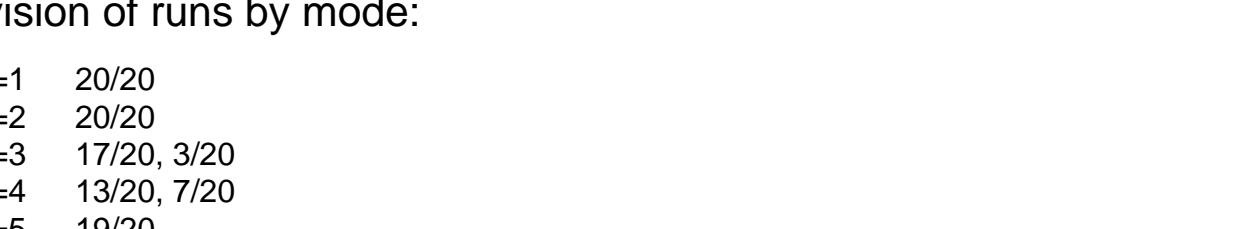
K=4 MinorCluster1



K=6 MinorCluster1



K=7 MinorCluster1



K=9 MinorCluster1



K=10 MinorCluster1



Division of runs by mode:

K=1 20/20

K=2 20/20

K=3 17/20, 3/20

K=4 13/20, 7/20

K=5 19/20

K=6 15/20, 4/20

K=7 12/20, 8/20

K=8 18/20

K=9 17/20, 2/20

K=10 13/20, 7/20

APPENDIX 4- 2: Results of STRUCTURE analysis for the 126 samples belonging to the Guinean group. Population assignment for K=2 and K=5

