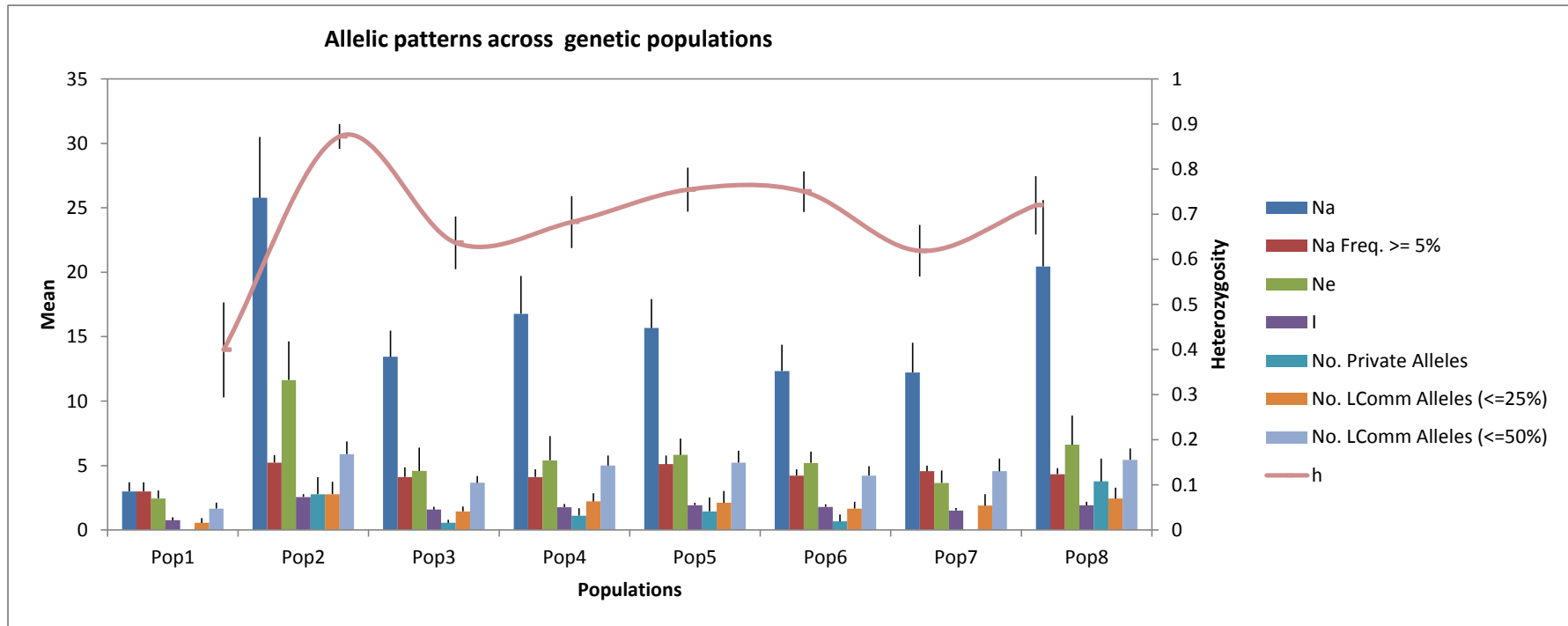


Mean values													
Population	Australia	Belgium	China	Cuba	France	Germany	India	Italy	Netherlands	Norway	Spain	Switzerland	USA
Na	3.778	17.333	4.000	6.667	14.000	16.333	11.444	3.444	32.778	19.222	19.000	17.111	16.222
Na Freq. >= 5%	3.778	5.556	4.000	6.667	5.778	5.222	5.556	3.444	5.333	4.444	6.111	5.111	7.222
Ne	3.381	8.075	3.378	5.158	7.659	8.232	8.155	3.259	10.563	6.745	9.111	9.117	9.621
I	1.205	2.352	1.141	1.709	2.183	2.265	2.153	1.194	2.551	2.142	2.389	2.367	2.353
No. Private Alleles	0.000	0.333	0.000	0.000	0.000	0.333	0.222	0.000	5.556	0.333	0.333	0.222	0.444
No. LComm Alleles (<=25%)	0.556	2.889	0.889	1.444	2.889	3.333	1.889	0.778	4.444	3.556	4.000	3.556	3.222
No. LComm Alleles (<=50%)	1.111	5.333	1.444	2.556	5.000	5.444	3.333	1.111	6.889	6.000	6.333	5.889	5.111
h	0.649	0.860	0.601	0.769	0.837	0.832	0.840	0.681	0.872	0.800	0.858	0.860	0.862
uh	0.811	0.872	0.687	0.854	0.854	0.848	0.875	0.907	0.873	0.805	0.867	0.874	0.877

**Standard Error (SE) values**

Population	Australia	Belgium	China	Cuba	France	Germany	India	Italy	Netherlands	Norway	Spain	Switzerland	USA
Na	0.364	1.675	0.707	0.553	1.732	2.392	1.365	0.176	5.883	2.900	2.255	2.163	2.332
Na Freq. >= 5%	0.364	0.338	0.707	0.553	0.401	0.434	0.530	0.176	0.408	0.294	0.588	0.309	0.722
Ne	0.461	1.282	0.758	0.640	1.627	1.628	1.308	0.234	2.775	1.657	2.162	1.776	2.125
I	0.130	0.111	0.184	0.131	0.146	0.198	0.167	0.061	0.188	0.192	0.160	0.163	0.174
No. Private Alleles	0.000	0.167	0.000	0.000	0.000	0.236	0.222	0.000	2.167	0.167	0.167	0.147	0.338
No. LComm Alleles (<=25%)	0.242	0.790	0.455	0.556	0.484	0.799	0.455	0.222	1.107	0.852	1.000	0.801	0.862
No. LComm Alleles (<=50%)	0.351	1.093	0.530	0.689	0.816	1.156	0.707	0.200	1.457	1.179	1.333	1.136	1.124
h	0.054	0.014	0.066	0.042	0.022	0.033	0.035	0.022	0.019	0.031	0.020	0.022	0.022
uh	0.068	0.014	0.076	0.047	0.023	0.034	0.037	0.029	0.019	0.031	0.021	0.022	0.023



**Mean values**

<b>Population</b>	<b>Pop1</b>	<b>Pop2</b>	<b>Pop3</b>	<b>Pop4</b>	<b>Pop5</b>	<b>Pop6</b>	<b>Pop7</b>	<b>Pop8</b>
<b>Na</b>	3.000	25.778	13.444	16.778	15.667	12.333	12.222	20.444
<b>Na Freq. &gt;= 5%</b>	3.000	5.222	4.111	4.111	5.111	4.222	4.556	4.333
<b>Ne</b>	2.445	11.634	4.579	5.399	5.828	5.202	3.650	6.616
<b>I</b>	0.756	2.558	1.581	1.770	1.924	1.796	1.515	1.909
<b>No. Private Alleles</b>	0.000	2.778	0.556	1.111	1.444	0.667	0.000	3.778
<b>No. LComm Alleles (&lt;=25%)</b>	0.556	2.778	1.444	2.222	2.111	1.667	1.889	2.444
<b>No. LComm Alleles (&lt;=50%)</b>	1.667	5.889	3.667	5.000	5.222	4.222	4.556	5.444
<b>h</b>	0.399	0.872	0.637	0.683	0.754	0.750	0.619	0.720
<b>uh</b>	0.456	0.876	0.644	0.687	0.758	0.759	0.624	0.722

**Standard Error (SE) values**

<b>Population</b>	<b>Pop1</b>	<b>Pop2</b>	<b>Pop3</b>	<b>Pop4</b>	<b>Pop5</b>	<b>Pop6</b>	<b>Pop7</b>	<b>Pop8</b>
<b>Na</b>	0.707	4.725	2.015	2.938	2.242	2.041	2.296	5.140
<b>Na Freq. &gt;= 5%</b>	0.707	0.572	0.754	0.588	0.676	0.494	0.444	0.471
<b>Ne</b>	0.625	3.004	1.813	1.883	1.265	0.894	0.969	2.259
<b>I</b>	0.227	0.215	0.209	0.250	0.189	0.204	0.201	0.280
<b>No. Private Alleles</b>	0.000	1.310	0.242	0.588	1.082	0.553	0.000	1.762
<b>No. LComm Alleles (&lt;=25%)</b>	0.377	0.969	0.377	0.619	0.920	0.527	0.904	0.835
<b>No. LComm Alleles (&lt;=50%)</b>	0.471	0.992	0.527	0.782	0.940	0.722	0.973	0.884
<b>h</b>	0.105	0.027	0.059	0.058	0.049	0.045	0.057	0.065
<b>uh</b>	0.120	0.028	0.059	0.058	0.049	0.046	0.058	0.065