

Tabela 1. Quantidade de Variantes Encontradas nas Amostras (VCF's)

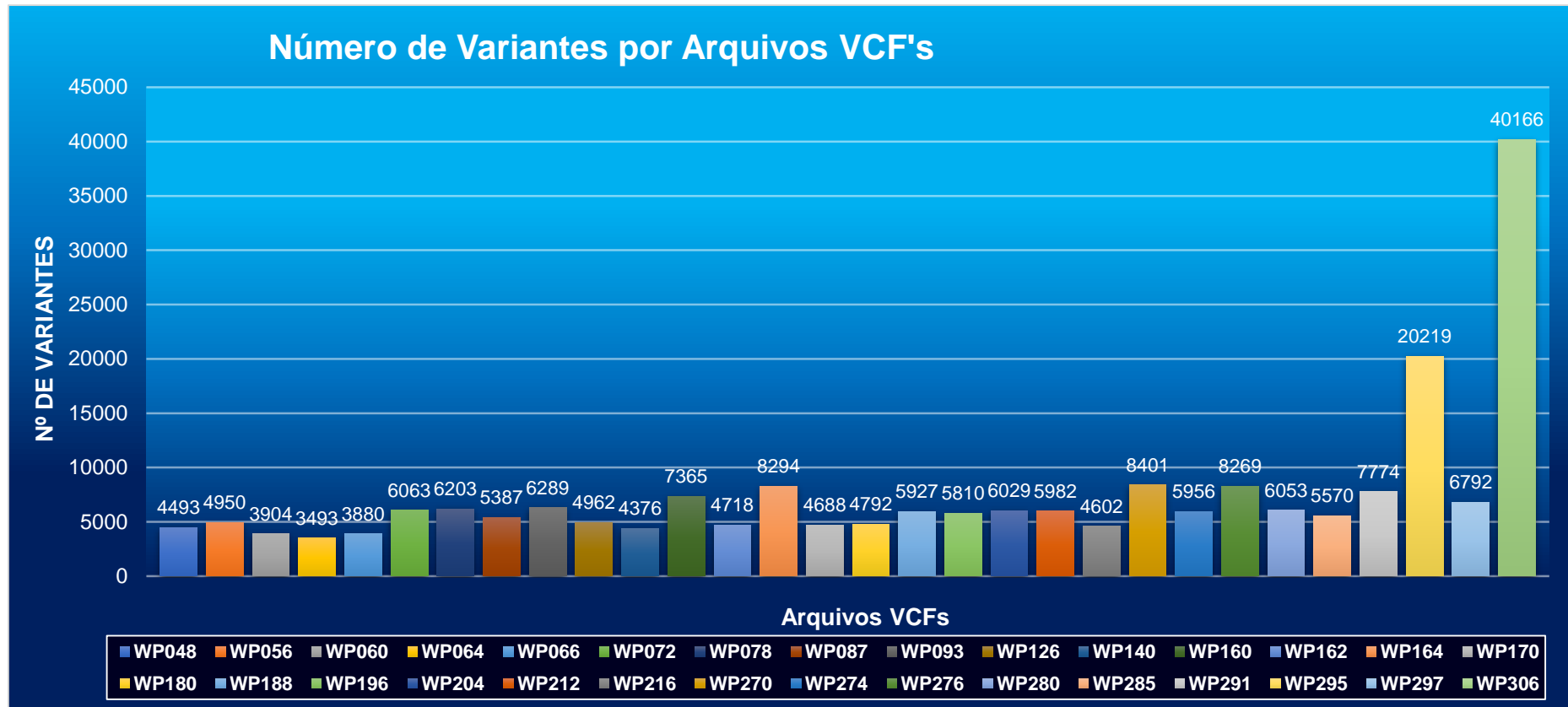


Tabela 2. Quantidade de Cada Classe de Variantes (Todas Amostras Juntas)

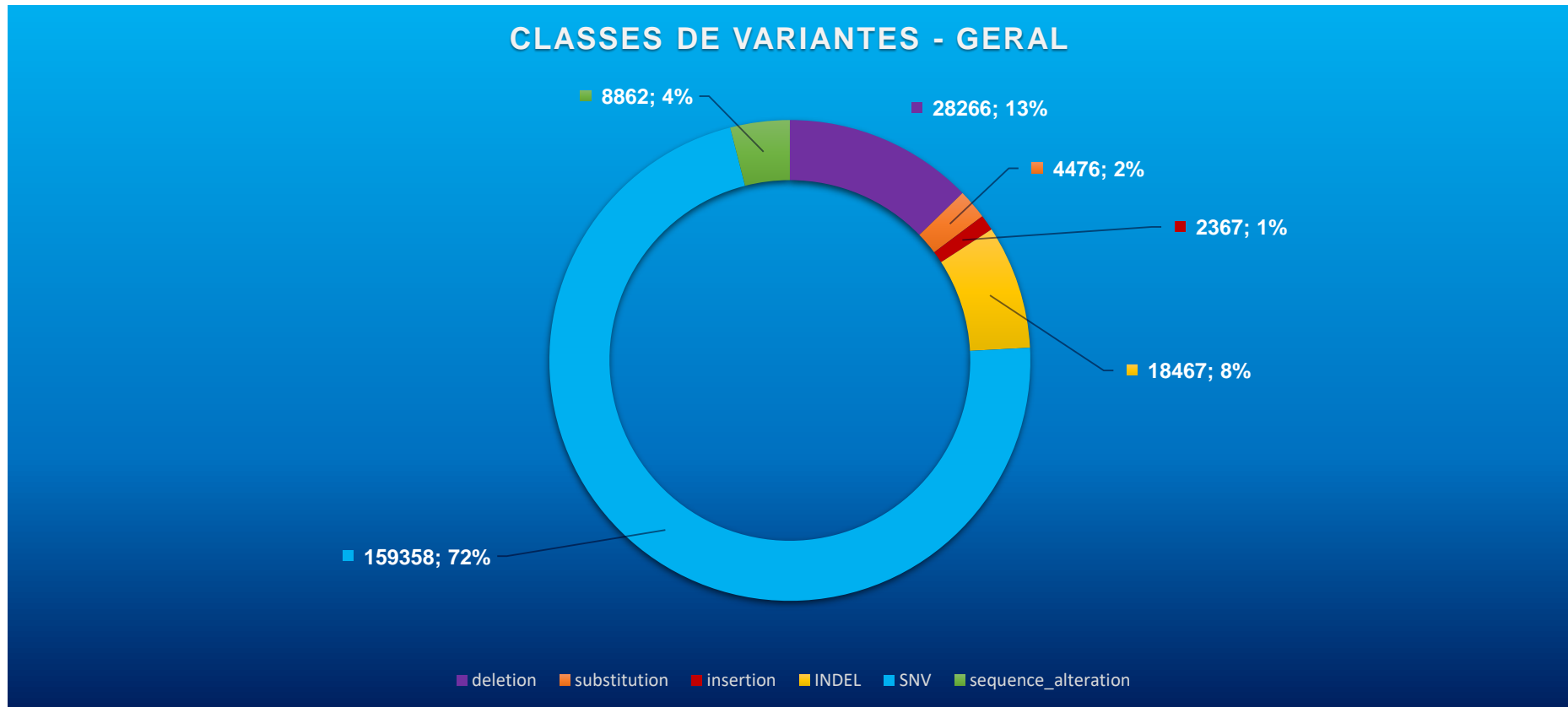
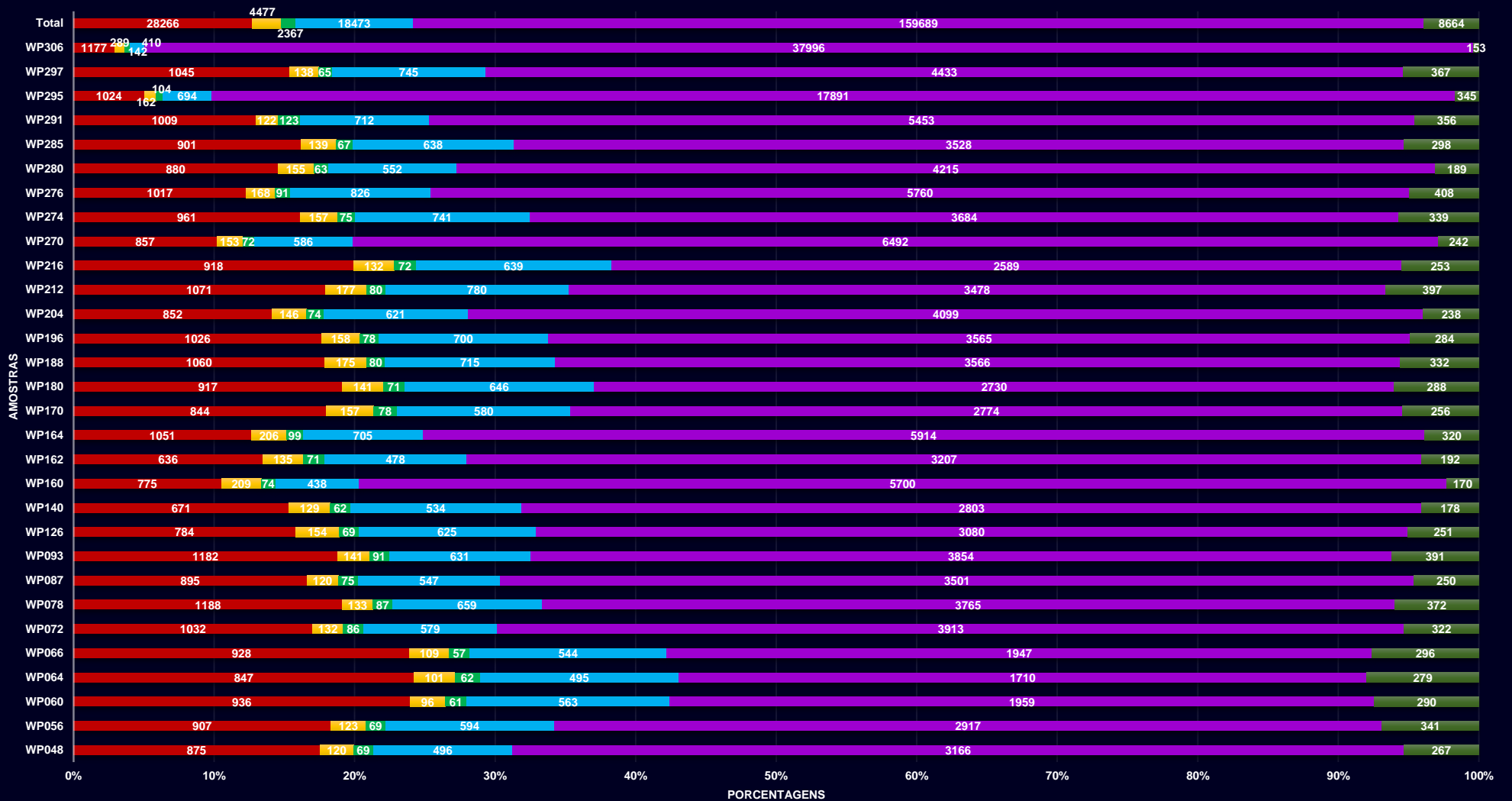


Tabela 3. Quantidade de Cada Classe de Variantes por Amostras

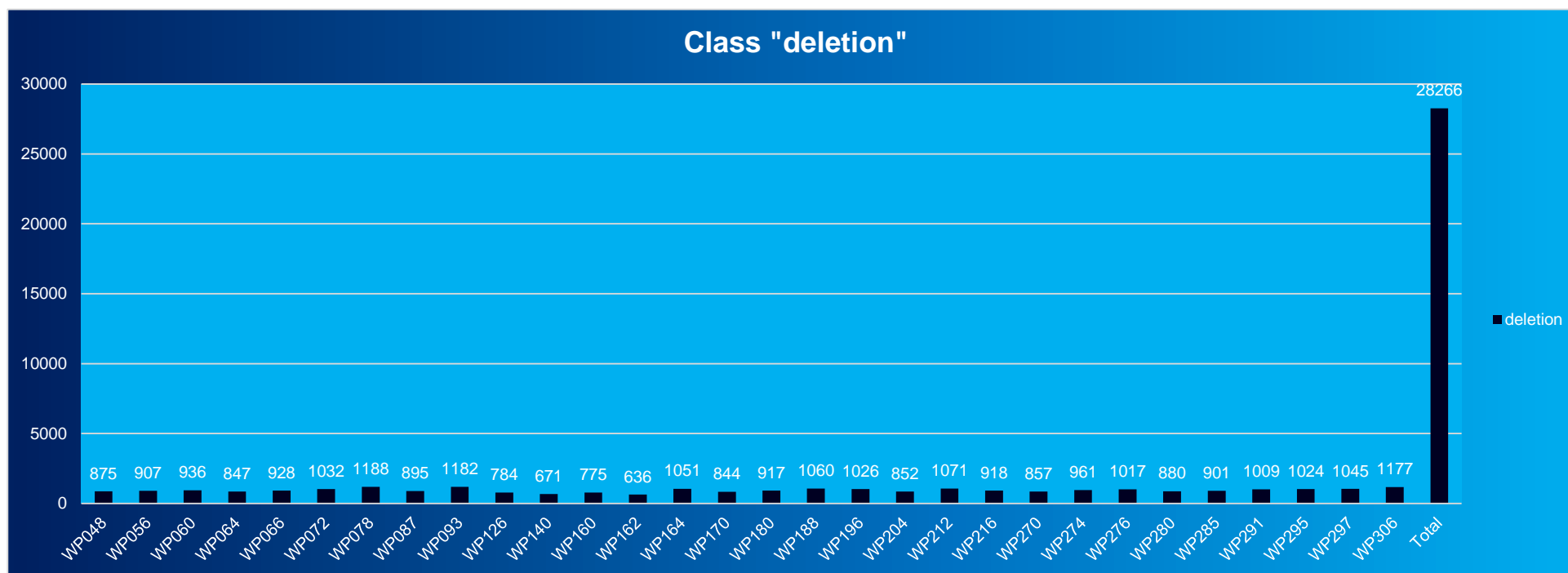
	deletion	substitution	insertion	INDEL	SNV	sequence_alteration
WP048	875	120	69	496	3166	267
WP056	907	123	69	594	2917	341
WP060	936	96	61	563	1959	290
WP064	847	101	62	495	1710	279
WP066	928	109	57	544	1947	296
WP072	1032	132	86	579	3913	322
WP078	1188	133	87	659	3765	372
WP087	895	120	75	547	3501	250
WP093	1182	141	91	631	3854	391
WP126	784	154	69	625	3080	251
WP140	671	129	62	534	2803	178
WP160	775	209	74	438	5700	170
WP162	636	135	71	478	3207	192
WP164	1051	206	99	705	5914	320
WP170	844	157	78	580	2774	256
WP180	917	141	71	646	2730	288
WP188	1060	175	80	715	3566	332
WP196	1026	158	78	700	3565	284
WP204	852	146	74	621	4099	238
WP212	1071	177	80	780	3478	397
WP216	918	132	72	639	2589	253
WP270	857	153	72	586	6492	242
WP274	961	157	75	741	3684	339
WP276	1017	168	91	826	5760	408
WP280	880	155	63	552	4215	189
WP285	901	139	67	638	3528	298
WP291	1009	122	123	712	5453	356
WP295	1024	162	104	694	17891	345
WP297	1045	138	65	745	4433	367
WP306	1177	289	142	410	37996	153
Total	28266	4477	2367	18473	159689	8664

Proporções de Classes de Variantes por Amostras

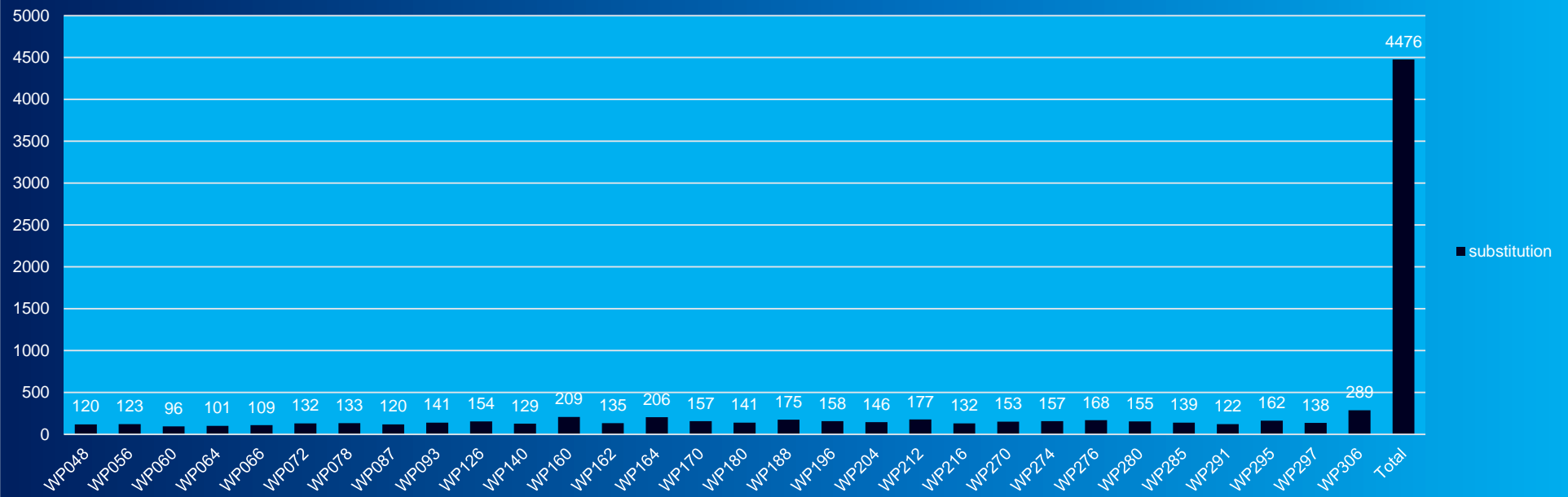


deletion substitution insertion INDEL SNV sequence_alteration

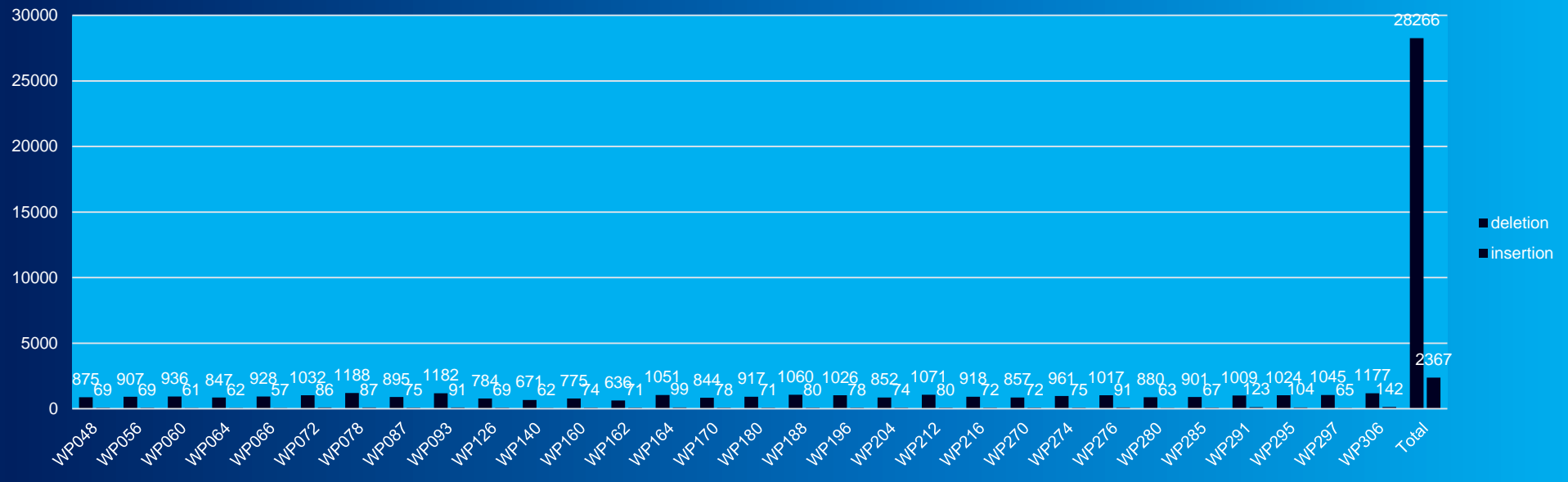
Tabela 4-9. Quantidade de Cada Classe de Variantes (Visão Individualizada)



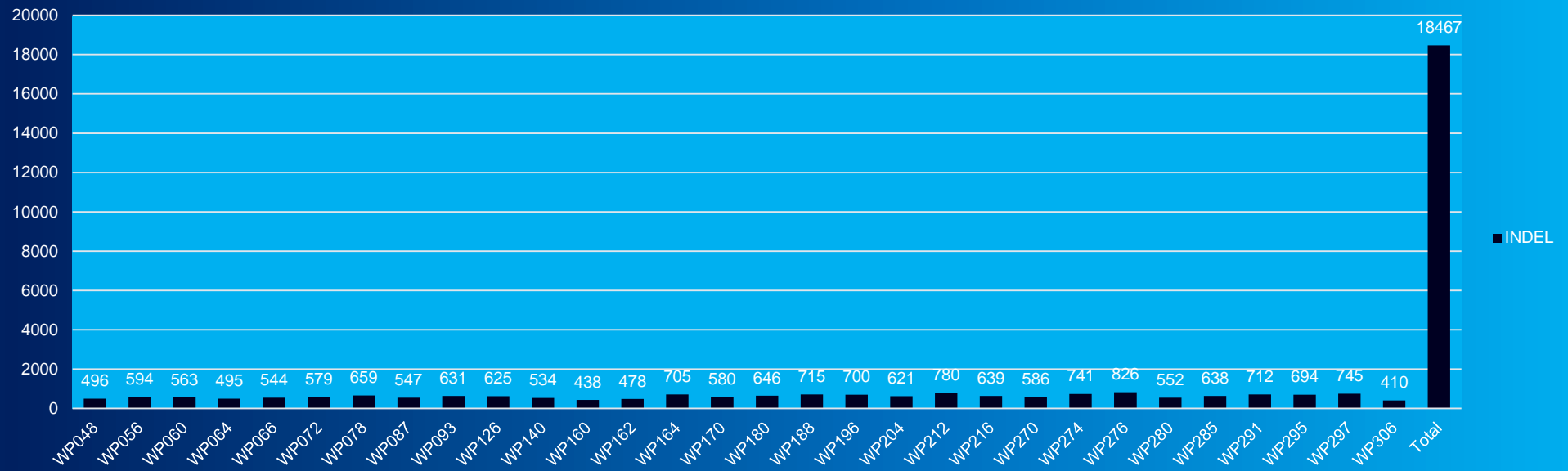
Class "substitution"



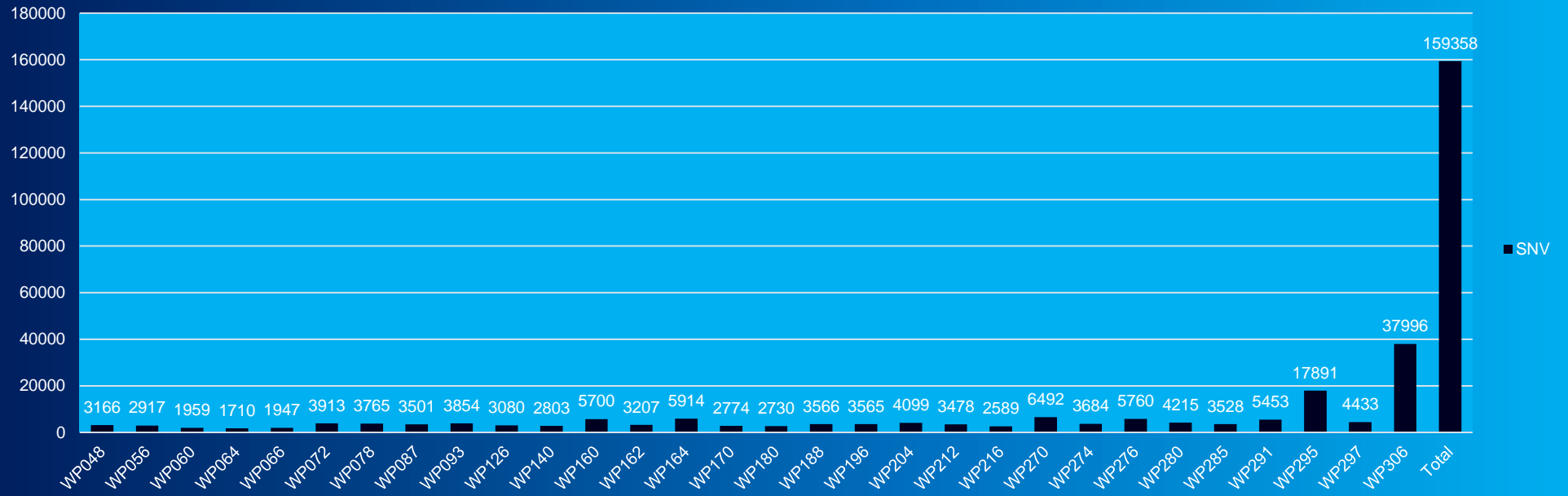
Class "insertion"



Class "INDEL "



Class "SNV"



Class "sequence_alteration"

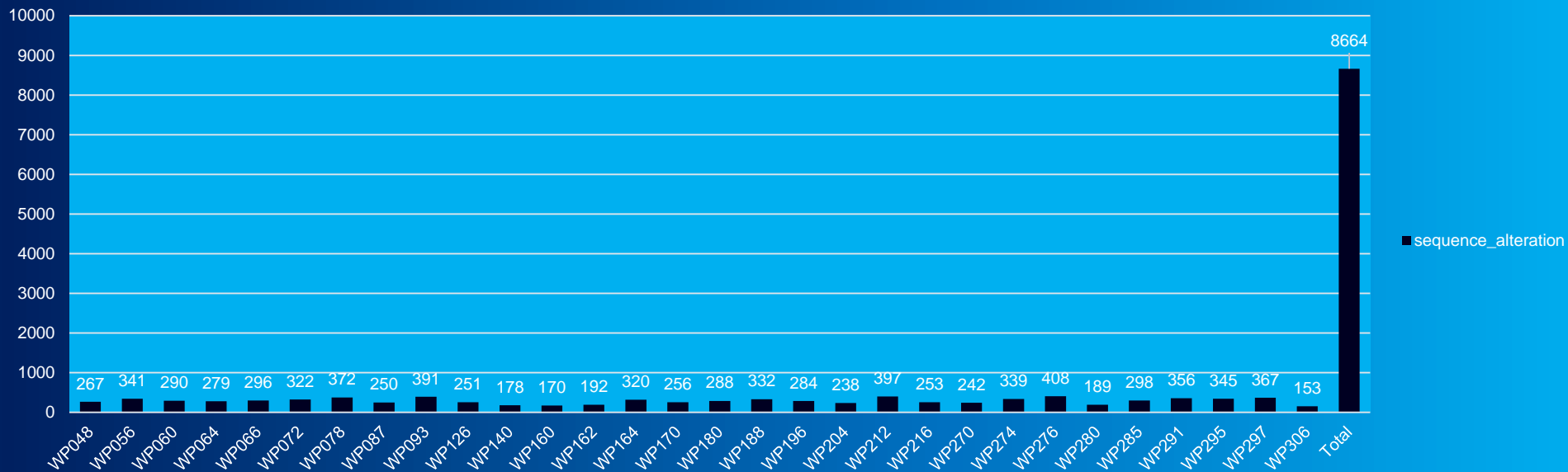


Tabela 10. Quantidades de Cada Tipo de Fator de Patogenicidade

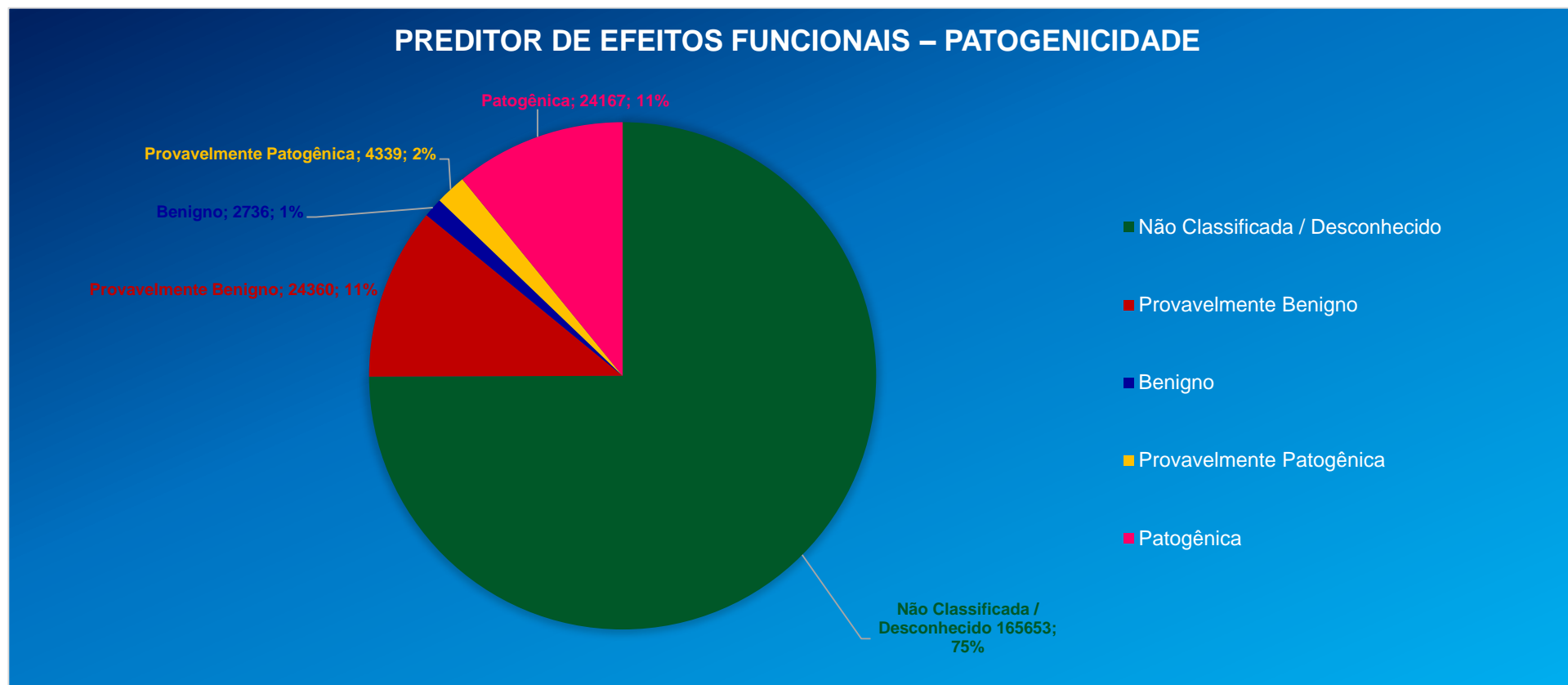


Tabela 11. Consequências Patogênicas

