

Table S1. List of 113 PI-associated mutations included in one or more of four expert 2020 lists: prevalence in PI-naïve and PI-experienced persons before and after 2009.

Pos ¹	AA ¹	Lists		<2009											≥2009												
				2009	2020	%PI (22,266)	%PI-Naive										%PI (10,534)	%PI-Naive									
		ALL (61,067)	A (4,316)				B (33,132)	C (8,935)	01 (5,265)	02 (3,743)	D (1,497)	F (726)	G (2,036)	Other (1,438)	ALL (54,813)	A (3,209)		B (21,435)	C (10,194)	01 (10,189)	02 (3,616)	D (891)	F (794)	G (1,425)	Other (3,039)		
10	C	1	1	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	
10	F	4	4	8.6	0.2	0.1	0.3	0.1	0.1	0.0	0.3	0.0	0.1	0.5	6.7	0.2	0.2	0.3	0.1	0.2	0.1	0.5	0.0	0.0	0.0	0.0	
10	I	4	3	31	8.5	14	8.9	2.3	10	6.0	6.1	9.1	11	26	16	8.4	20	9.2	2.2	7.7	7.7	6.6	10	19	15	15	
10	M	1	1	0.2	0.3	0.6	0.0	0.8	0.3	0.1	0.2	0.1	1.5	1.4	0.3	0.5	0.8	0.2	1.2	0.2	0.5	0.0	0.3	2.5	2.0	2.0	
10	R	3	1	0.4	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.2	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	
10	V	4	3	11	4.8	8.5	3.4	1.8	5.9	11	8.1	33	0.5	7.6	11	5.9	12	4.4	1.7	5.9	12	8.6	50	1.6	4.8	4.8	
11	I	4	4	3.0	0.6	0.5	0.3	0.5	0.5	3.3	0.5	0.0	1.1	1.4	1.4	0.8	1.1	0.3	0.5	0.6	4.6	0.2	0.6	1.6	0.3	0.3	
11	L	1	1	0.7	0.0	0.0	0.0	0.1	0.0	0.0	0.1	0.0	0.0	0.0	0.2	0.0	0.2	0.0	0.0	0.0	0.0	0.1	0.0	0.1	0.0	0.0	
15	A	1	1	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.1	0.1	
15	V	1	1	33	28	15	22	85	12	10	17	75	17	32	41	33	19	25	81	14	16	18	60	22	25	25	
16	E	3	1	6.2	9.4	20	4.2	7.2	25	23	9.0	26	4.5	16	13	14	24	6.2	9.1	26	31	9.8	30	6.0	19	19	
20	I	4	2	8.4	9.9	2.7	0.1	0.1	2.7	95	0.7	0.4	98	8.9	6.2	11	5.4	0.2	0.1	1.9	95	2.4	3.1	97	9.8	9.8	
20	M	4	3	3.3	0.5	0.1	0.4	0.6	0.1	0.6	1.3	1.3	0.9	1.8	1.7	0.5	0.1	0.6	0.6	0.0	0.5	2.7	2.3	0.6	1.1	1.1	
20	R	4	3	17	8.2	17	2.8	19	16	1.7	12	43	0.0	18	18	13	20	3.0	22	27	2.3	13	31	0.4	11	11	
20	T	4	3	5.3	0.1	0.1	0.1	0.1	0.1	0.1	0.2	0.0	0.0	0.8	4.1	0.1	0.0	0.1	0.1	0.2	0.1	0.1	0.0	0.0	0.0	0.0	
20	V	3	2	1.1	0.1	0.0	0.0	0.0	0.0	1.5	0.2	0.0	0.6	0.2	0.5	0.1	0.1	0.0	0.0	0.0	1.4	0.0	0.0	0.9	0.5	0.5	
23	I	2	2	1.3	0.0	0.2	0.0	0.0	0.0	0.0	0.1	0.0	0.0	0.0	1.1	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	
24	F	2	1	0.6	0.0	0.1	0.0	0.1	0.0	0.0	0.1	0.0	0.0	0.0	0.3	0.0	0.0	0.0	0.0	0.0	0.0	0.1	0.0	0.0	0.0	0.0	
24	I	4	4	5.6	0.0	0.0	0.0	0.0	0.0	0.0	0.1	0.0	0.0	0.0	2.5	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	
24	M	0	1	0.1	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.1	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	
30	N	4	4	6.2	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	2.4	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	
32	I	4	4	5.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	2.5	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.1	0.0	0.0	
32	ins	1	1	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	
33	F	4	4	12	0.4	1.0	0.2	0.1	1.4	0.4	0.7	0.0	0.4	0.5	6.9	0.5	1.6	0.1	0.2	1.2	0.6	0.4	0.3	0.2	0.7	0.7	
33	I	4	3	2.2	0.6	0.3	1.0	0.0	0.2	0.2	1.4	0.1	0.2	0.6	1.7	0.7	0.8	1.4	0.0	0.1	0.2	0.9	0.6	0.2	0.6	0.6	
33	V	4	2	1.8	1.7	0.5	2.6	0.1	0.5	0.2	7.0	0.6	0.3	0.8	1.4	1.3	0.7	2.6	0.3	0.3	0.4	6.4	1.0	0.1	0.8	0.8	
34	Q	1	1	2.6	0.1	0.0	0.0	0.0	0.0	0.5	0.0	0.0	0.7	0.0	1.3	0.1	0.0	0.0	0.0	0.0	0.7	0.0	0.0	0.4	0.3	0.3	
35	G	3	1	1.1	0.2	0.1	0.1	0.1	0.1	1.5	0.1	0.1	0.3	1.2	0.7	0.2	0.1	0.1	0.1	0.1	1.4	0.2	0.0	0.4	0.7	0.7	
35	N	1	1	1.6	0.6	1.0	0.1	0.2	1.7	2.6	0.3	0.7	2.9	0.3	1.9	1.0	3.0	0.1	0.3	1.1	3.8	1.9	0.9	3.1	1.5	1.5	
36	I	4	2	49	51	98	17	84	97	97	63	97	98	79	62	62	97	18	85	97	97	71	96	98	69	69	

36	L	3	2	2.6	1.3	0.4	1.0	3.6	0.3	0.9	2.5	0.7	0.3	1.8	2.5	1.4	0.6	1.2	3.6	0.2	0.8	1.3	0.8	0.5	1.0
36	V	3	2	2.1	1.0	0.4	0.4	4.7	0.6	0.2	0.5	0.1	0.2	1.2	1.4	1.1	0.2	0.4	3.4	0.4	0.3	0.6	0.8	0.1	2.7
38	W	1	1	0.2	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.1	0.0	0.0	0.1	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0
41	T	1	1	0.1	0.0	0.0	0.0	0.1	0.0	0.1	0.1	0.0	0.1	0.3	0.1	0.0	0.0	0.0	0.1	0.0	0.2	0.0	0.0	0.0	0.1
43	R	1	1	2.4	2.7	1.4	3.0	1.3	2.4	5.6	1.6	3.7	3.1	5.0	3.0	3.2	2.7	3.4	1.3	2.7	7.8	4.5	2.3	4.1	3.9
43	T	3	3	5.6	0.1	0.1	0.1	0.1	0.0	0.2	0.1	0.1	0.1	1.2	3.0	0.1	0.1	0.1	0.1	0.0	0.2	0.0	0.0	0.0	0.4
46	I	4	4	21	0.3	0.3	0.3	0.2	0.3	0.2	0.1	0.0	0.0	0.2	13	0.3	0.3	0.4	0.1	0.6	0.2	0.1	0.4	0.2	0.0
46	L	4	4	9.2	0.3	0.3	0.4	0.1	0.3	0.1	0.2	0.1	0.4	0.3	3.4	0.4	0.2	0.3	0.1	0.9	0.1	0.0	0.4	0.0	0.6
46	V	1	1	0.5	0.0	0.0	0.0	0.1	0.1	0.1	0.0	0.0	0.0	0.0	0.4	0.1	0.0	0.1	0.0	0.1	0.1	0.0	0.1	0.0	0.1
47	A	4	4	0.4	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	1.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0
47	V	4	4	4.5	0.0	0.0	0.0	0.0	0.0	0.1	0.0	0.0	0.1	0.0	2.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.1	0.0
48	A	2	2	0.5	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.2	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.1
48	L	0	1	0.1	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0
48	M	2	2	0.5	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.2	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0
48	Q	0	1	0.1	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.1	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0
48	S	1	1	0.1	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.1	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0
48	T	1	1	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0
48	V	4	4	3.7	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.7	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0
50	L	4	4	1.5	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	2.5	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0
50	V	4	4	1.8	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	1.2	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0
53	L	4	4	5.9	0.0	0.0	0.0	0.0	0.0	0.0	0.1	0.1	0.0	0.0	3.0	0.0	0.0	0.0	0.0	0.0	0.0	0.1	0.0	0.0	0.0
53	W	1	1	0.1	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0
53	Y	3	2	0.4	0.0	0.0	0.0	0.0	0.0	0.1	0.0	0.0	0.0	0.0	0.2	0.0	0.0	0.0	0.0	0.0	0.2	0.1	0.0	0.0	0.0
54	A	4	4	1.3	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.4	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0
54	L	4	4	3.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	1.4	0.0	0.1	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0
54	M	4	4	2.3	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.1	0.0	0.0	0.6	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.1
54	S	4	3	0.7	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.1	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0
54	T	4	4	0.8	0.0	0.1	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.2	0.2	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.1	0.0	0.0
54	V	4	4	23	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	12	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0
58	E	4	3	6.3	0.3	0.1	0.4	0.4	0.0	0.0	1.2	0.1	0.1	0.2	4.3	0.4	0.1	0.6	0.5	0.0	0.0	0.7	0.5	0.3	0.4
60	E	2	2	12	9.0	5.2	8.9	17	2.8	0.6	16	11	2.1	37	13	15	6.2	18	19	2.9	1.0	15	27	2.4	32
62	V	3	3	34	17	6.3	27	4.6	4.7	1.4	19	19	3.2	20	25	17	7.9	33	5.1	5.5	2.2	21	23	6.3	16
63	P	3	2	63	40	15	55	36	12	19	35	9.1	20	14	49	37	17	54	37	17	22	39	9.6	24	33
64	L	1	1	2.9	3.6	1.2	4.5	1.9	0.6	7.3	3.1	2.6	3.6	2.4	3.4	3.6	2.3	4.7	2.2	0.6	9.8	5.2	2.8	5.1	2.9
64	M	2	2	1.7	2.6	2.0	1.6	1.6	0.8	13	0.8	2.1	9.6	3.6	2.2	3.3	3.4	1.9	1.9	0.6	18	0.9	1.9	13	2.1
64	V	2	3	18	11	0.8	16	0.5	0.4	1.7	66	0.6	0.4	13	13	7.1	1.2	14	0.8	0.6	1.8	57	1.5	0.7	5.5
66	V	0	1	1.1	0.1	0.1	0.1	0.1	0.0	0.4	0.1	0.0	0.1	1.4	0.4	0.1	0.1	0.0	0.1	0.0	0.2	0.0	0.1	0.4	0.4

69	I	1	1	0.1	0.0	0.0	0.0	0.0	0.1	0.1	0.1	0.0	0.0	0.0	0.1	0.0	0.1	0.0	0.0	0.1	0.1	0.0	0.0	0.1	0.0
69	K	3	2	15	41	96	1.5	98	97	96	2.8	1.4	94	61	41	54	92	2.6	97	96	96	6.4	3.7	89	58
69	N	1	1	1.0	0.5	0.1	0.7	0.1	0.1	0.2	0.3	0.1	0.3	0.2	0.2	0.3	0.3	0.6	0.0	0.2	0.3	0.1	0.6	0.4	0.4
69	Q	1	1	3.3	2.4	2.1	2.9	1.1	0.7	1.0	5.1	8.4	1.4	11	4.2	2.8	4.4	4.0	1.3	0.7	1.1	16	7.3	1.1	6.7
69	R	1	2	1.5	0.6	0.7	0.4	0.3	1.0	1.2	0.3	1.0	4.0	1.2	1.0	0.9	1.3	0.4	0.4	0.9	2.3	0.3	0.6	8.4	0.8
69	Y	1	1	2.2	1.9	0.0	2.5	0.0	0.0	0.0	18	7.2	0.0	1.2	1.9	1.4	0.2	2.3	0.0	0.0	0.0	9.7	17.9	0.0	0.7
71	I	4	2	3.2	0.1	0.0	0.1	0.0	0.0	0.0	0.0	0.0	0.0	0.0	1.7	0.1	0.0	0.3	0.0	0.0	0.0	0.0	0.0	0.0	0.0
71	L	3	2	0.4	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.2	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0
71	T	4	3	9.9	4.7	0.1	8.2	1.0	0.2	0.2	1.9	0.1	0.4	0.3	6.3	4.7	0.2	11	1.0	0.3	0.3	3.0	0.5	0.5	2.5
71	V	4	2	27	3.3	0.0	5.9	0.2	0.3	0.2	0.3	0.1	0.3	0.5	12	3.5	0.2	8.0	0.3	0.2	0.1	0.6	0.1	1.1	6.1
73	A	4	2	0.6	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.2	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0
73	C	3	3	1.1	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.3	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0
73	D	0	1	0.1	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.1	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.1	0.1	0.0
73	S	4	4	8.3	0.0	0.0	0.0	0.0	0.0	0.1	0.0	0.0	0.0	0.3	2.7	0.0	0.0	0.0	0.0	0.0	0.1	0.0	0.0	0.0	0.2
73	T	4	4	2.4	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.6	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0
73	V	1	1	0.2	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.1	0.0	0.0	0.1	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0
74	A	1	1	1.8	0.6	0.3	0.5	2.2	0.2	0.1	0.0	1.1	0.3	0.3	1.4	0.8	0.4	0.6	2.5	0.3	0.2	0.1	0.5	0.1	0.2
74	P	3	4	1.9	0.1	0.0	0.1	0.0	0.1	0.0	0.1	0.0	0.0	0.2	1.3	0.0	0.0	0.0	0.0	0.0	0.0	0.1	0.0	0.0	0.0
74	S	2	1	7.3	2.3	1.5	0.4	11	0.9	1.3	0.3	7.9	0.8	1.4	7.1	3.0	2.5	0.3	12	1.0	1.2	0.1	3.0	0.6	1.5
76	V	4	4	3.3	0.0	0.0	0.0	0.0	0.0	0.0	0.1	0.1	0.0	0.0	3.3	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0
77	I	3	2	27	21	12	33	7.4	2.0	2.6	15	10	2.7	26	26	19	12	37	7.4	4.0	3.3	13	15	2.5	21
82	A	4	4	22	0.0	0.0	0.1	0.0	0.0	0.0	0.0	0.1	0.0	0.0	12	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.1	0.1	0.0
82	C	3	2	0.6	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.4	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0
82	F	4	4	1.8	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	1.1	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0
82	G	1	1	0.1	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.1	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0
82	L	3	3	0.3	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.2	0.0	0.0	0.1	0.0	0.0	0.0	0.0	0.0	0.0	0.0
82	M	3	2	0.3	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.4	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.1	0.0	0.0
82	S	4	4	1.3	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.5	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0
82	T	4	4	2.8	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.8	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0
83	D	2	3	0.9	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.6	0.0	0.0	0.0	0.0	0.1	0.0	0.0	0.0	0.1	0.0
84	A	3	3	0.1	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0
84	C	2	2	0.1	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0
84	V	4	4	12	0.0	0.0	0.0	0.0	0.0	0.0	0.1	0.0	0.0	0.0	3.8	0.0	0.0	0.1	0.0	0.0	0.0	0.0	0.0	0.0	0.1
85	V	3	3	4.6	0.2	0.1	0.2	0.1	0.1	0.1	0.5	0.0	0.0	0.2	2.1	0.2	0.4	0.2	0.1	0.1	0.1	0.2	0.0	0.0	0.2
88	D	4	4	5.4	0.0	0.0	0.0	0.0	0.1	0.0	0.0	0.0	0.0	0.0	2.7	0.0	0.1	0.0	0.0	0.0	0.0	0.0	0.0	0.1	0.0
88	G	1	1	0.2	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.1	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0
88	S	4	4	1.6	0.0	0.0	0.0	0.0	0.0	0.1	0.0	0.0	0.1	0.0	1.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0

88	T	1	1	0.1	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.1	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0
89	I	3	2	2.9	1.3	1.6	0.1	2.3	2.1	5.9	0.2	1.0	2.4	11	4.0	2.0	2.6	0.2	3.2	2.1	7.9	0.1	1.4	3.6	5.8
89	M	2	1	14	40	97	1.6	82	96	93	2.1	80	96	55	38	51	95	1.9	76	94	92	4.9	83	95	51
89	R	1	1	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.1	0.0	0.1
89	T	3	2	0.4	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.9	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0
89	V	4	4	3.7	0.1	0.0	0.0	0.1	0.0	0.4	0.1	0.1	0.1	0.2	1.4	0.0	0.0	0.0	0.0	0.1	0.2	0.0	0.1	0.1	0.1
90	M	4	4	30	0.2	0.0	0.3	0.2	0.1	0.1	0.1	0.1	0.3	0.2	9.6	0.2	0.1	0.4	0.1	0.0	0.2	0.0	0.1	0.4	0.1
93	L	2	1	40	41	21	42	96	19	3.5	3.9	8.8	4.2	21	45	49	17	53	96	35	0.7	11	14	2.5	42
93	M	3	1	0.7	0.1	0.3	0.1	0.0	0.2	0.3	0.1	0.6	0.2	0.6	0.2	0.1	0.3	0.1	0.0	0.3	0.4	0.0	0.4	0.0	0.4

Abbreviation: Pos – amino acid position; AA – amino acid; List – Number of expert lists with the mutation in 2009 and 2020; %PI – percent prevalence among PI-experienced persons; %PI-Naïve – percent prevalence among PI-naïve persons.

The parenthetical numbers in the headers represent the number of persons from whom a sequence was available.

¹40 PI-associated surveillance drug resistance mutations at 18 positions (Bennett 2009) were indicated in bold.

Table S2. List of 66 NRTI-associated mutations included in one or more of four expert 2020 lists: prevalence in RTI-naïve and NRTI-experienced persons before and after 2009.

[illegible]

Abbreviation: Pos – amino acid position; AA – amino acid; List – Number of expert lists with the mutation in 2009 and 2020; %NRTI – percent prevalence among NRTI-experienced persons; %RTI-Naïve – percent prevalence among RTI-naïve persons.

The parenthetical numbers in the headers represent the number of persons from whom a sequence was available

¹34 NRTI-associated surveillance drug resistance mutations at 15 positions (Bennett 2009) were indicated in bold.

Table S3. List of 69 NNRTI-associated mutations included in one or more of four expert 2020 lists: prevalence in RTI-naïve and NNRTI-experienced persons before and after 2009.

Pos ¹	AA ¹	Lists		<2009											≥2009												
				%NNRTI	%RTI-Naïve										%NNRTI	%RTI-Naïve											
		2009	2020		(27,755)	ALL (54,910)	A (4,245)	B (30,428)	C (7,596)	01 (4,571)	02 (2,871)	D (1,279)	F (606)	G (1,003)		Other (2,361)	(22,219)	ALL (45,940)	A (2,423)	B (18,847)	C (4,811)	01 (9,495)	02 (2,382)	D (612)	F (777)	G (495)	Other (6,048)
90	I	2	3	5.9	1.9	1.4	2.1	1.0	0.5	6.8	2.6	0.2	1.2	0.8	8.6	1.6	1.9	1.7	1.1	0.5	7.2	1.6	0.0	3.6	0.5		
98	G	4	4	6.4	0.2	0.1	0.1	0.8	0.2	0.1	0.1	0.5	0.3	0.0	8.3	0.2	0.1	0.2	0.6	0.1	0.3	0.2	0.9	1.0	0.2		
100	I	4	4	4.5	0.0	0.0	0.0	0.1	0.0	0.0	0.2	0.0	0.0	0.0	3.5	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0		
100	V	1	1	0.1	0.0	0.0	0.1	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.1	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.1		
101	E	4	4	7.2	0.1	0.1	0.2	0.1	0.0	0.1	0.4	0.0	0.1	0.0	9.3	0.2	0.2	0.2	0.3	0.1	0.1	0.2	0.0	0.8	0.1		
101	H	1	3	1.3	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	1.3	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0		
101	I	0	1	0.1	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.2	0.0	0.0	0.1	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0		
101	N	2	1	0.5	0.0	0.0	0.0	0.0	0.0	0.1	0.0	0.0	0.0	0.0	0.7	0.0	0.1	0.0	0.0	0.0	0.0	0.0	0.2	0.0	0.0		
101	P	4	4	1.7	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	1.4	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0		
101	Q	2	1	4.9	0.5	0.4	0.6	0.5	0.3	0.6	0.2	0.2	0.7	0.1	3.9	0.7	0.9	0.9	0.6	0.4	0.8	0.5	0.5	0.2	0.9		
101	R	2	2	1.6	1.0	0.3	1.2	0.9	0.7	0.3	1.1	0.8	0.6	2.0	1.6	1.1	0.7	1.2	1.1	0.9	0.6	0.5	0.6	0.0	1.0		
101	T	0	1	0.4	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.2	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0		
103	H	2	3	0.1	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0		
103	N	4	4	36	1.3	0.5	1.8	1.1	0.2	0.7	0.9	1.5	0.9	0.7	36	1.6	2.1	2.3	1.5	0.4	2.4	2.3	1.5	3.0	1.3		
103	R	2	2	2.4	1.9	0.2	2.9	1.2	0.8	0.2	0.4	0.3	0.2	0.4	2.8	2.0	0.3	3.1	1.2	2.1	0.7	0.3	1.0	0.6	1.6		
103	S	3	4	2.0	0.0	0.0	0.0	0.1	0.0	0.0	0.0	0.2	0.0	0.0	2.3	0.1	0.2	0.1	0.0	0.0	0.0	0.2	0.0	0.0	0.1		
103	T	3	3	0.2	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.1	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0		
106	A	4	4	1.8	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.1	0.0	1.4	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.1		
106	I	4	4	3.9	1.9	1.3	2.2	0.0	4.8	0.5	1.6	4.6	1.1	2.0	3.5	3.0	1.7	4.5	0.2	2.9	1.4	2.1	16	2.2	1.1		
106	M	4	4	2.6	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	9.1	0.1	0.0	0.0	0.2	0.1	0.0	0.0	0.0	0.2	0.0		
106	T	0	1	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0		
108	I	3	3	8.6	0.5	0.7	0.5	0.2	0.2	0.7	0.3	0.3	0.1	0.4	9.4	0.4	0.3	0.5	0.2	0.2	0.9	0.3	0.3	0.2	0.1		
138	A	0	4	3.1	2.5	3.2	1.9	6.0	0.4	2.7	3.0	3.3	2.2	1.6	4.9	2.2	6.2	1.9	5.4	0.3	2.7	2.1	1.7	1.8	0.6		
138	G	0	4	0.7	0.2	0.2	0.2	0.3	0.4	0.1	0.5	2.0	0.1	0.1	1.1	0.4	0.2	0.3	0.3	0.2	0.3	0.3	0.5	0.2	1.6		
138	K	2	4	0.4	0.1	0.0	0.1	0.2	0.0	0.1	0.3	0.0	0.0	0.0	0.7	0.1	0.0	0.1	0.1	0.1	0.1	0.3	0.1	0.2	0.2		
138	Q	0	4	1.0	0.0	0.1	0.0	0.0	0.0	0.0	0.1	0.2	0.0	0.0	1.6	0.1	0.1	0.1	0.0	0.0	0.0	0.0	0.0	0.2	0.0		
138	R	0	4	0.0	0.0	0.0	0.0	0.0	0.0	0.1	0.0	0.0	0.0	0.0	0.1	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0		
138	S	0	2	0.0	0.0	0.0	0.0	0.3	0.0	0.0	0.0	0.0	0.0	0.0	0.1	0.0	0.1	0.0	0.1	0.0	0.0	0.0	0.0	0.0	0.0		
179	D	4	4	2.2	1.6	0.2	2.1	1.1	2.2	0.0	0.2	3.8	0.0	0.7	3.3	2.4	0.5	3.1	1.8	3.3	0.0	0.5	0.8	0.0	1.9		
179	E	2	2	1.2	0.6	0.0	0.6	0.1	0.2	0.8	0.4	0.0	4.3	0.5	1.8	1.5	0.0	1.4	0.3	0.7	1.0	0.2	0.3	3.4	15		

179	F	4	3	0.3	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.1	0.2	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0
179	I	2	1	13	8.9	54	4.3	3.1	16.2	3.9	7.0	5.3	0.4	6.6	16	10	51	5.4	3.7	20	5.3	6.9	17	0.8	6.1
179	L	0	4	0.2	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.3	0.0	0.0	0.0	0.0	0.0	0.1	0.0	0.0	0.0	0.0
179	M	1	1	0.1	0.0	0.0	0.0	0.0	0.0	0.2	0.0	0.0	0.0	0.0	0.3	0.0	0.0	0.0	0.0	0.0	0.1	0.0	0.1	0.0	0.0
179	T	1	3	0.5	0.3	1.6	0.1	0.0	1.1	0.1	1.0	0.2	0.0	0.4	0.9	0.8	1.7	0.2	0.1	2.9	0.2	1.5	0.1	0.0	0.9
181	C	4	4	18	0.2	0.1	0.1	0.3	0.2	0.1	0.2	0.0	0.2	0.0	18	0.2	0.2	0.1	0.2	0.1	0.3	0.7	0.1	0.4	0.3
181	F	0	1	0.2	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.2	0.0	0.1	0.0	0.1	0.0	0.0	0.0	0.0	0.0	0.0
181	G	0	1	0.1	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.1	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0
181	I	4	4	0.8	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.6	0.0	0.1	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0
181	S	1	1	0.1	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.1	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.1	0.0	0.0
181	V	4	4	0.5	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.7	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0
188	C	4	4	0.7	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	1.0	0.0	0.1	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0
188	F	1	2	0.6	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.6	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.1
188	H	4	4	0.8	0.0	0.0	0.0	0.0	0.0	0.0	0.1	0.0	0.0	0.1	0.7	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0
188	L	4	4	4.5	0.0	0.0	0.1	0.0	0.0	0.0	0.0	0.0	0.3	0.0	4.4	0.1	0.0	0.1	0.0	0.0	0.0	0.5	0.0	0.0	0.1
189	I	0	1	2.6	1.2	0.9	1.2	0.6	0.7	3.3	1.8	0.8	2.7	0.7	2.3	1.3	1.2	1.3	0.9	1.3	2.5	1.6	1.0	1.8	1.0
190	A	4	4	15	0.2	0.0	0.2	0.1	0.1	0.1	0.2	0.0	0.0	0.0	15	0.2	0.3	0.2	0.3	0.1	0.1	4.7	0.1	0.0	0.1
190	C	3	3	0.1	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.1	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0
190	E	3	4	0.5	0.0	0.0	0.0	0.0	0.1	0.0	0.0	0.0	0.0	0.0	0.6	0.1	0.0	0.0	0.1	0.1	0.0	0.0	0.0	0.0	0.1
190	Q	3	3	0.3	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.4	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0
190	S	4	4	2.7	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	3.3	0.0	0.0	0.0	0.0	0.0	0.0	0.2	0.0	0.0	0.0
190	T	3	3	0.1	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.1	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0
190	V	3	3	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.1	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0
221	Y	1	4	6.5	0.1	0.1	0.1	0.1	0.0	0.1	0.2	0.2	0.0	0.1	8.1	0.1	0.1	0.1	0.1	0.1	0.0	0.2	0.0	0.4	0.1
225	H	4	4	4.3	0.0	0.0	0.0	0.0	0.0	0.1	0.0	0.0	0.0	0.0	7.4	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0
227	C	2	4	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.1	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0
227	I	0	2	0.0	0.0	0.0	0.1	0.0	0.0	0.1	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0
227	L	2	3	2.2	0.1	0.0	0.1	0.0	0.1	0.0	0.0	0.2	0.0	0.0	3.5	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0
227	R	0	1	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0
227	V	0	1	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.2	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0
230	I	0	4	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.1	0.0	0.1	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0
230	L	3	4	1.5	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	2.6	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.1
230	V	0	1	0.1	0.0	0.0	0.0	0.0	0.1	0.1	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0
234	I	1	3	0.3	0.0	0.1	0.0	0.0	0.0	0.1	0.1	0.0	0.0	0.0	0.4	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.1
236	L	2	2	0.1	0.0	0.0	0.0	0.0	0.0	0.2	0.0	0.2	0.0	0.2	0.2	0.0	0.0	0.1	0.0	0.0	0.0	0.0	0.0	0.0	0.0
238	N	1	2	0.6	0.0	0.0	0.1	0.0	0.0	0.1	0.1	0.2	0.1	0.0	0.4	0.1	0.0	0.1	0.0	0.0	0.2	0.0	0.0	0.0	0.0
238	T	2	2	2.7	0.1	0.1	0.1	0.0	0.0	0.0	0.0	0.2	0.0	0.0	3.0	0.0	0.0	0.1	0.0	0.0	0.1	0.2	0.0	0.0	0.0

318	F	2	2	1.2	0.1	0.1	0.1	0.1	0.0	0.2	0.0	0.8	0.0	0.0	1.3	0.2	0.0	0.1	0.2	0.1	0.8	0.0	0.0	1.9	0.0
348	I	0	1	14	0.2	0.3	0.2	0.4	0.0	0.2	0.0	0.0	0.0	0.0	7.0	0.0	0.0	0.0	0.0	0.1	0.0	0.0	0.0	0.0	0.0

Abbreviation: Pos – amino acid position; AA – amino acid; List – Number of expert lists with the mutation in 2009 and 2020; %NNRTI – percent prevalence among NNRTI-experienced persons; %RTI-Naïve – percent prevalence among RTI-naïve persons.
The parenthetical numbers in the headers represent the number of persons from whom a sequence was available
¹19 NNRTI-associated surveillance drug resistance mutations at 10 positions (Bennett 2009) were indicated in bold.

Table S4. Number of sequences according to treatment and subtype

Subtype	#Protease sequences		#RT sequences	
	PI-Naïve (n=115,880)	PI-Experienced (n=32,800)	RTI-Naïve (n=100,850)	RTI-Experienced (n=56,573)
All	115,880	328,00	100,850	56,573
A	7,525	1,532	6,668	3,873
B	54,567	22,236	49,275	29,132
C	19,129	3,617	12,407	12,109
CRF01_AE	15,454	696	14,066	3,061
CRF02_AG	7,359	366	5,253	2,289
D	2,388	390	1,891	983
F	1,520	1,529	1,383	1,611
G	3461	401	1,498	1,548
CRF06_cpx	393	7	1,134	356
CRF07_BC	651	9	3,311	16
CRF08_BC	225	29	814	29
CRF12_BF	411	1,164	267	673
CRF18_cpx	224	40	104	55
CRF35_AD	430	30	241	16
CRF45_cpx	320	49	64	48
CRF63_02A1	128	77	127	86
Other	1,695	628	2,347	688

Table S6. NRTI-associated SDRMs and candidate SDRMs: Comparisons of prevalence in RTI-naïve and NRTI-experienced persons before and after 2009 using Fisher's exact test.

Pos	AA	SDRM	RTI-Naïve					NRTI-Experienced				
			% in <2009 (n=54,910)	% in ≥2009 (n=45,940)	OR	p	Adjusted p	% in <2009 (n=32,363)	% in ≥2009 (n=24,210)	OR	p	Adjusted p
41	L	SDRM	0.003	0.002	0.62	5.9E-05	2.0E-03	0.330	0.148	0.35	0.0E+00	0.0E+00
65	R	SDRM	0.000	0.000	0.85	4.3E-01	1.0E+00	0.040	0.104	2.80	3.0E-198	8.9E-197
67	E	SDRM	0.000	0.000	0.75	3.0E-01	1.0E+00	0.006	0.002	0.37	1.1E-11	1.2E-10
67	G	SDRM	0.000	0.000	0.50	6.9E-02	1.0E+00	0.024	0.023	0.93	9.3E-02	3.7E-01
67	N	SDRM	0.001	0.000	0.52	2.7E-02	6.9E-01	0.302	0.157	0.43	0.0E+00	0.0E+00
69	D	SDRM	0.001	0.001	1.17	2.8E-01	1.0E+00	0.069	0.031	0.43	8.9E-95	2.3E-93
69	ins	SDRM	0.000	0.000	Inf	4.6E-01	1.0E+00	0.010	0.002	0.21	1.5E-33	2.9E-32
70	E	SDRM	0.000	0.000	0.64	2.4E-01	1.0E+00	0.006	0.026	4.04	3.6E-79	9.0E-78
70	R	SDRM	0.001	0.000	0.52	8.3E-03	2.5E-01	0.206	0.124	0.54	1.6E-151	4.6E-150
74	I	SDRM	0.000	0.000	1.02	5.5E-01	1.0E+00	0.046	0.037	0.79	2.6E-08	2.6E-07
74	V	SDRM	0.000	0.000	0.40	2.1E-01	1.0E+00	0.086	0.039	0.43	5.0E-115	1.4E-113
75	A	SDRM	0.000	0.000	0.60	3.5E-01	1.0E+00	0.008	0.002	0.32	4.4E-18	6.5E-17
75	M	SDRM	0.000	0.001	1.62	8.7E-02	1.0E+00	0.036	0.035	0.96	2.0E-01	3.7E-01
75	S	SDRM	0.000	0.000	0.00	5.4E-01	1.0E+00	0.003	0.001	0.47	8.3E-05	5.0E-04
75	T	SDRM	0.000	0.000	0.00	1.0E+00	1.0E+00	0.015	0.005	0.32	2.4E-33	4.1E-32
77	L	SDRM	0.002	0.001	0.42	1.3E-05	4.6E-04	0.019	0.012	0.62	6.8E-12	8.1E-11
115	F	SDRM	0.000	0.000	0.20	9.7E-02	1.0E+00	0.020	0.045	2.27	3.0E-62	7.3E-61
116	Y	SDRM	0.000	0.000	0.00	8.8E-02	1.0E+00	0.022	0.010	0.43	1.4E-32	2.3E-31
151	M	SDRM	0.000	0.000	0.00	1.0E+00	1.0E+00	0.029	0.012	0.43	4.8E-42	9.6E-41
184	I	SDRM	0.000	0.001	1.91	3.3E-02	8.2E-01	0.023	0.033	1.45	4.5E-13	5.8E-12
184	V	SDRM	0.002	0.001	0.70	1.0E-02	3.0E-01	0.545	0.606	1.28	9.2E-48	2.0E-46
210	W	SDRM	0.001	0.001	1.55	3.5E-02	8.5E-01	0.208	0.077	0.32	0.0E+00	0.0E+00
215	C	SDRM	0.001	0.001	0.92	3.9E-01	1.0E+00	0.009	0.004	0.42	8.3E-15	1.2E-13

215	D	SDRM	0.003	0.002	0.64	6.4E-04	2.1E-02	0.008	0.004	0.55	4.7E-08	3.7E-07
215	E	SDRM	0.001	0.002	1.25	1.1E-01	1.0E+00	0.002	0.001	0.75	1.2E-01	3.7E-01
215	F	SDRM	0.000	0.000	0.60	5.7E-01	1.0E+00	0.118	0.083	0.68	3.9E-42	8.2E-41
215	I	SDRM	0.000	0.000	1.05	5.1E-01	1.0E+00	0.022	0.020	0.94	1.4E-01	3.7E-01
215	S	SDRM	0.004	0.003	0.82	4.5E-02	1.0E+00	0.025	0.018	0.73	3.2E-08	2.9E-07
215	V	SDRM	0.000	0.000	0.80	5.8E-01	1.0E+00	0.009	0.005	0.57	6.8E-08	4.8E-07
215	Y	SDRM	0.000	0.000	0.09	2.6E-03	7.9E-02	0.305	0.134	0.35	0.0E+00	0.0E+00
219	E	SDRM	0.000	0.000	0.94	5.2E-01	1.0E+00	0.068	0.061	0.90	7.5E-04	3.7E-03
219	N	SDRM	0.000	0.000	0.34	1.1E-02	3.0E-01	0.034	0.014	0.42	4.5E-50	1.0E-48
219	Q	SDRM	0.001	0.001	0.72	7.8E-02	1.0E+00	0.126	0.069	0.51	5.9E-113	1.6E-111
219	R	SDRM	0.001	0.000	0.57	2.6E-02	6.9E-01	0.027	0.013	0.47	2.3E-33	4.1E-32
44	A	Candidate	0.000	0.000	3.59	2.5E-01	1.0E+00	0.017	0.006	0.33	1.8E-37	3.1E-36
65	N	Candidate	0.000	0.000	0.28	2.5E-02	4.2E-01	0.001	0.002	1.77	1.1E-02	5.2E-02
67	H	Candidate	0.000	0.000	3.59	2.5E-01	1.0E+00	0.002	0.001	0.55	2.8E-03	2.5E-02
67	S	Candidate	0.000	0.000	0.00	3.0E-01	1.0E+00	0.003	0.002	0.58	8.3E-04	9.1E-03
67	T	Candidate	0.000	0.000	0.00	1.0E+00	1.0E+00	0.002	0.002	0.63	1.0E-02	5.2E-02
69	A	Candidate	0.002	0.003	1.24	4.8E-02	7.1E-01	0.011	0.009	0.79	3.6E-03	2.9E-02
69	del	Candidate	0.000	0.000	0.00	5.4E-01	1.0E+00	0.001	0.003	2.30	9.4E-06	1.1E-04
69	G	Candidate	0.000	0.000	0.00	5.4E-01	1.0E+00	0.003	0.001	0.56	1.7E-03	1.7E-02
70	G	Candidate	0.000	0.000	0.00	1.0E+00	1.0E+00	0.005	0.003	0.72	8.6E-03	5.1E-02
70	N	Candidate	0.000	0.000	0.96	5.2E-01	1.0E+00	0.003	0.004	1.31	3.4E-02	1.0E-01
70	Q	Candidate	0.000	0.000	0.22	2.5E-02	4.2E-01	0.002	0.006	2.76	8.1E-13	1.3E-11
70	T	Candidate	0.000	0.000	1.05	5.1E-01	1.0E+00	0.003	0.007	2.25	1.6E-11	2.2E-10
215	A	Candidate	0.001	0.001	0.80	1.4E-01	1.0E+00	0.003	0.003	0.91	3.0E-01	4.6E-01
215	L	Candidate	0.001	0.001	1.05	4.8E-01	1.0E+00	0.002	0.002	0.84	2.3E-01	4.6E-01
215	N	Candidate	0.000	0.000	0.96	5.6E-01	1.0E+00	0.013	0.007	0.55	2.3E-12	3.4E-11
219	H	Candidate	0.000	0.000	0.00	1.0E+00	1.0E+00	0.003	0.001	0.41	8.4E-07	1.1E-05
219	W	Candidate	0.000	0.000	0.00	1.0E+00	1.0E+00	0.003	0.002	0.60	3.9E-03	2.9E-02

Estimated odds ratio (OR) by Fisher's exact test, their p-values and adjusted p values by Holm's method are shown. Significant increase or decrease in prevalence with adjusted p <0.05 is indicated in bold.

Table S7. NNRTI-associated SDRMs and candidate SDRMs: Comparisons of prevalence in RTI-naïve and NNRTI-experienced persons before and after 2009 using Fisher's exact test.

Pos	AA	SDRM	RTI-Naive					NNRTI-Experienced				
			% in <2009 (n=54,910)	% in ≥2009 (n=45,940)	OR	p	Adjusted p	% in <2009 (n=27,755)	% in ≥2009 (n=22,219)	OR	p	Adjusted p
100	I	SDRM	0.000	0.000	0.15	2.2E-03	3.5E-02	0.045	0.035	0.77	5.2E-09	7.7E-08
101	E	SDRM	0.001	0.002	1.35	3.2E-02	4.2E-01	0.072	0.093	1.34	5.3E-19	9.0E-18
101	P	SDRM	0.000	0.000	1.20	7.9E-01	1.0E+00	0.017	0.014	0.81	1.8E-03	1.8E-02
103	N	SDRM	0.013	0.016	1.24	2.2E-05	4.2E-04	0.358	0.356	0.99	3.4E-01	1.0E+00
103	S	SDRM	0.000	0.001	1.87	3.4E-02	4.2E-01	0.020	0.023	1.16	7.6E-03	6.1E-02
106	A	SDRM	0.000	0.000	0.51	1.2E-01	9.6E-01	0.018	0.014	0.76	7.1E-05	9.2E-04
106	M	SDRM	0.000	0.001	3.93	5.3E-04	9.5E-03	0.026	0.091	3.75	6.0E-224	1.1E-222
179	F	SDRM	0.000	0.000	0.30	9.4E-02	8.4E-01	0.003	0.002	0.76	8.7E-02	5.7E-01
181	C	SDRM	0.002	0.002	1.05	4.2E-01	1.0E+00	0.179	0.181	1.01	3.3E-01	1.0E+00
181	I	SDRM	0.000	0.000	2.39	2.6E-01	1.0E+00	0.008	0.006	0.73	2.4E-03	2.2E-02
181	V	SDRM	0.000	0.000	Inf	4.6E-01	1.0E+00	0.005	0.007	1.43	1.2E-03	1.3E-02
188	C	SDRM	0.000	0.000	0.80	3.9E-01	1.0E+00	0.007	0.010	1.44	2.0E-04	2.4E-03
188	H	SDRM	0.000	0.000	0.32	2.5E-02	3.5E-01	0.008	0.007	0.88	1.4E-01	6.9E-01
188	L	SDRM	0.000	0.001	1.56	5.5E-02	5.7E-01	0.045	0.044	0.98	3.7E-01	1.0E+00
190	A	SDRM	0.002	0.002	1.54	1.5E-03	2.6E-02	0.145	0.146	1.01	4.0E-01	1.0E+00
190	E	SDRM	0.000	0.001	2.72	3.3E-03	4.9E-02	0.005	0.006	1.20	8.2E-02	5.7E-01
190	S	SDRM	0.000	0.000	2.39	1.2E-01	9.6E-01	0.027	0.033	1.24	2.1E-05	2.9E-04
225	H	SDRM	0.000	0.000	0.36	5.1E-02	5.7E-01	0.043	0.074	1.79	6.1E-51	1.1E-49
230	L	SDRM	0.000	0.000	1.17	4.9E-01	1.0E+00	0.015	0.026	1.67	9.7E-16	1.6E-14
101	H	Candidate	0.000	0.000	Inf	4.3E-02	6.5E-01	0.013	0.013	1.00	5.2E-01	1.0E+00
101	N	Candidate	0.000	0.000	1.20	4.2E-01	1.0E+00	0.005	0.007	1.31	1.3E-02	1.0E-01
101	T	Candidate	0.000	0.000	1.59	2.7E-01	1.0E+00	0.004	0.002	0.64	5.3E-03	5.8E-02

[illegible]

Table S8. PI-associated SDRMs and candidate SDRMs: Comparisons of prevalence in PI-naïve and PI-experienced persons before and after 2009 using Fisher's exact test.

Pos	AA	SDRM	PII-Naive					PI-Experienced				
			% in <2009 (n=61,067)	% in ≥2009 (n=54,813)	OR	p	Adjusted p	% in <2009 (n=22,266)	% in ≥2009 (n=10,534)	OR	p	Adjusted p
23	I	SDRM	0.000	0.000	1.00	5.7E-01	1.0E+00	0.013	0.011	0.80	2.1E-02	8.5E-02
24	I	SDRM	0.000	0.000	0.26	1.9E-02	6.9E-01	0.056	0.025	0.43	2.6E-40	8.2E-39
30	N	SDRM	0.000	0.000	1.25	4.1E-01	1.0E+00	0.062	0.024	0.38	6.4E-54	2.0E-52
32	I	SDRM	0.000	0.000	0.37	1.0E-01	1.0E+00	0.050	0.025	0.49	1.2E-27	2.7E-26
46	I	SDRM	0.003	0.003	1.31	8.0E-03	3.0E-01	0.212	0.126	0.54	2.9E-82	9.7E-81
46	L	SDRM	0.003	0.004	1.20	4.2E-02	1.0E+00	0.092	0.034	0.35	4.5E-88	1.6E-86
47	A	SDRM	0.000	0.000	0.00	1.0E+00	1.0E+00	0.004	0.010	2.43	7.8E-10	1.2E-08
47	V	SDRM	0.000	0.000	0.77	3.5E-01	1.0E+00	0.045	0.020	0.42	4.7E-34	1.3E-32
48	M	SDRM	0.000	0.000	0.00	1.0E+00	1.0E+00	0.005	0.002	0.28	2.2E-08	3.0E-07
48	V	SDRM	0.000	0.000	0.00	1.0E+00	1.0E+00	0.037	0.007	0.18	1.0E-67	3.4E-66
50	L	SDRM	0.000	0.000	0.00	4.1E-02	1.0E+00	0.015	0.025	1.70	1.9E-10	3.0E-09
50	V	SDRM	0.000	0.000	0.28	7.6E-02	1.0E+00	0.018	0.012	0.66	1.9E-05	1.9E-04
53	L	SDRM	0.000	0.000	0.72	2.3E-01	1.0E+00	0.059	0.030	0.49	7.3E-32	1.8E-30
53	Y	SDRM	0.000	0.000	2.23	3.4E-02	1.0E+00	0.004	0.002	0.35	9.9E-06	1.1E-04
54	A	SDRM	0.000	0.000	0.00	1.0E+00	1.0E+00	0.013	0.004	0.27	1.1E-18	2.4E-17
54	L	SDRM	0.000	0.000	0.56	2.5E-01	1.0E+00	0.030	0.014	0.45	3.4E-21	7.6E-20
54	M	SDRM	0.000	0.000	1.11	7.1E-01	1.0E+00	0.023	0.006	0.24	1.5E-35	4.4E-34
54	S	SDRM	0.000	0.000	0.00	1.0E+00	1.0E+00	0.007	0.001	0.21	1.1E-12	1.9E-11
54	T	SDRM	0.000	0.000	0.64	3.4E-01	1.0E+00	0.008	0.002	0.19	7.1E-16	1.4E-14
54	V	SDRM	0.000	0.000	1.67	3.1E-01	1.0E+00	0.231	0.120	0.46	1.2E-131	4.4E-130
73	A	SDRM	0.000	0.000	1.11	6.5E-01	1.0E+00	0.006	0.002	0.34	8.1E-08	1.1E-06
73	C	SDRM	0.000	0.000	1.67	4.5E-01	1.0E+00	0.011	0.003	0.30	7.5E-15	1.4E-13
73	S	SDRM	0.000	0.000	1.67	1.1E-01	1.0E+00	0.083	0.027	0.31	4.6E-92	1.6E-90

73	T	SDRM	0.000	0.000	0.00	1.0E+00	1.0E+00	0.024	0.006	0.25	6.0E-35	1.7E-33
76	V	SDRM	0.000	0.000	1.34	4.3E-01	1.0E+00	0.033	0.033	1.00	4.9E-01	4.9E-01
82	A	SDRM	0.000	0.000	0.96	5.1E-01	1.0E+00	0.220	0.122	0.49	2.7E-106	1.0E-104
82	C	SDRM	0.000	0.000	Inf	2.2E-01	1.0E+00	0.006	0.004	0.66	8.7E-03	4.3E-02
82	F	SDRM	0.000	0.000	12.25	1.8E-03	7.0E-02	0.018	0.011	0.59	1.9E-07	2.3E-06
82	L	SDRM	0.000	0.000	1.35	2.5E-01	1.0E+00	0.003	0.002	0.69	7.6E-02	2.3E-01
82	M	SDRM	0.000	0.000	Inf	2.2E-01	1.0E+00	0.003	0.004	1.28	1.2E-01	2.4E-01
82	S	SDRM	0.000	0.000	0.56	5.4E-01	1.0E+00	0.013	0.005	0.40	6.0E-12	1.0E-10
82	T	SDRM	0.000	0.000	0.00	1.0E+00	1.0E+00	0.028	0.008	0.27	8.2E-38	2.5E-36
83	D	SDRM	0.000	0.000	1.62	1.5E-01	1.0E+00	0.009	0.006	0.69	4.1E-03	2.5E-02
84	A	SDRM	0.000	0.000	0.00	1.0E+00	1.0E+00	0.001	0.000	0.08	2.8E-04	1.9E-03
84	C	SDRM	0.000	0.000	0.00	1.0E+00	1.0E+00	0.001	0.000	0.00	6.2E-05	5.0E-04
84	V	SDRM	0.000	0.000	1.98	7.0E-02	1.0E+00	0.122	0.038	0.28	6.8E-151	2.7E-149
85	V	SDRM	0.002	0.002	0.95	4.1E-01	1.0E+00	0.046	0.021	0.44	6.9E-33	1.8E-31
88	D	SDRM	0.000	0.000	0.67	2.3E-01	1.0E+00	0.054	0.027	0.50	3.4E-29	8.0E-28
88	S	SDRM	0.000	0.000	0.00	1.3E-04	5.1E-03	0.016	0.010	0.64	2.1E-05	1.9E-04
90	M	SDRM	0.002	0.002	0.97	4.2E-01	1.0E+00	0.296	0.096	0.25	0.0E+00	0.0E+00
10	F	Candidate	0.002	0.002	1.01	4.9E-01	1.0E+00	0.086	0.067	0.76	1.5E-09	2.0E-08
10	R	Candidate	0.000	0.000	0.74	3.8E-01	1.0E+00	0.004	0.002	0.59	1.1E-02	4.2E-02
11	L	Candidate	0.000	0.000	0.67	1.4E-01	1.0E+00	0.007	0.002	0.36	2.7E-07	2.7E-06
20	T	Candidate	0.001	0.001	0.91	3.4E-01	1.0E+00	0.053	0.041	0.77	1.8E-06	1.6E-05
24	F	Candidate	0.000	0.000	0.15	2.1E-05	3.6E-04	0.006	0.003	0.47	5.0E-05	3.0E-04
34	Q	Candidate	0.001	0.001	1.01	5.2E-01	1.0E+00	0.026	0.013	0.49	1.8E-15	2.7E-14
41	T	Candidate	0.000	0.000	1.11	4.0E-01	1.0E+00	0.001	0.001	1.27	3.2E-01	5.4E-01
43	T	Candidate	0.001	0.001	0.92	3.4E-01	1.0E+00	0.056	0.030	0.52	3.9E-27	6.6E-26
46	V	Candidate	0.000	0.001	1.44	9.4E-02	1.0E+00	0.005	0.004	0.84	1.8E-01	5.4E-01
48	A	Candidate	0.000	0.000	2.23	4.6E-01	1.0E+00	0.005	0.002	0.36	9.4E-06	7.5E-05
66	V	Candidate	0.001	0.001	0.87	2.8E-01	1.0E+00	0.011	0.004	0.40	4.5E-10	6.3E-09
71	I	Candidate	0.001	0.001	1.34	7.6E-02	1.0E+00	0.032	0.017	0.51	4.4E-17	7.0E-16
71	L	Candidate	0.000	0.000	0.00	2.8E-01	1.0E+00	0.004	0.002	0.58	7.7E-03	3.8E-02

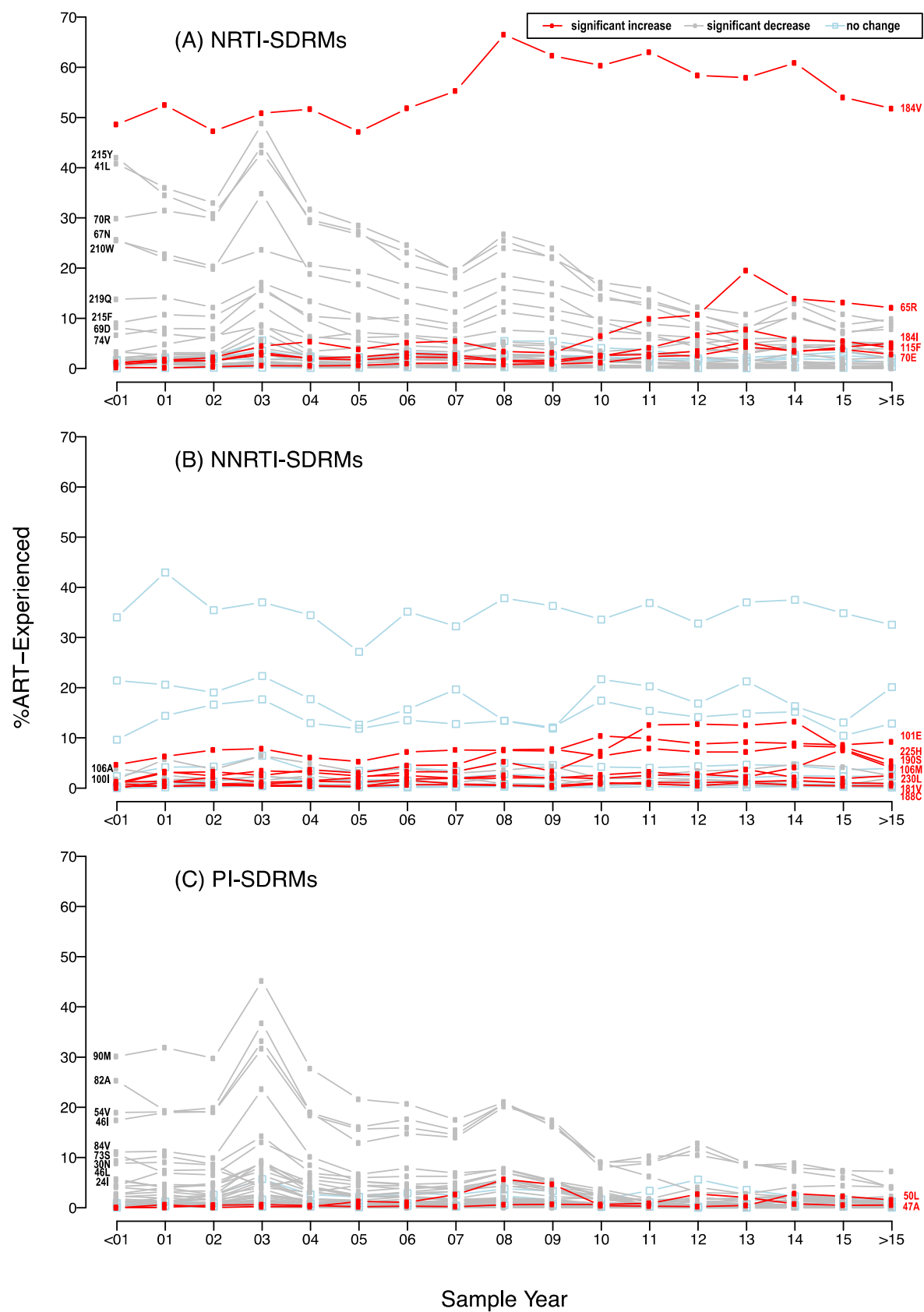


Figure S1. Yearly prevalence of (A) NRTI-SDRMs in NRTI-experienced persons, (B) NNRTI-SDRMs in NNRTI-experienced persons and (C) PI-SDRMs in PI-experienced persons. SDRMs significantly increased in prevalence are shown in red and indicated in red font on the right of the plotted lines. SDRMs significantly decreased in prevalence are shown in grey and indicated in black font on the left of the plotted lines. SDRMs did not change significantly in prevalence are shown in light blue. For SDRMs that decreased with a very low overall prevalence or did not change in prevalence, the labels are not shown. Full lists of NRTI-SDRMs, NNRTI-SDRMs and PI-SDRMs according to the change in prevalence are shown in Table 1, 3, and 5, respectively.

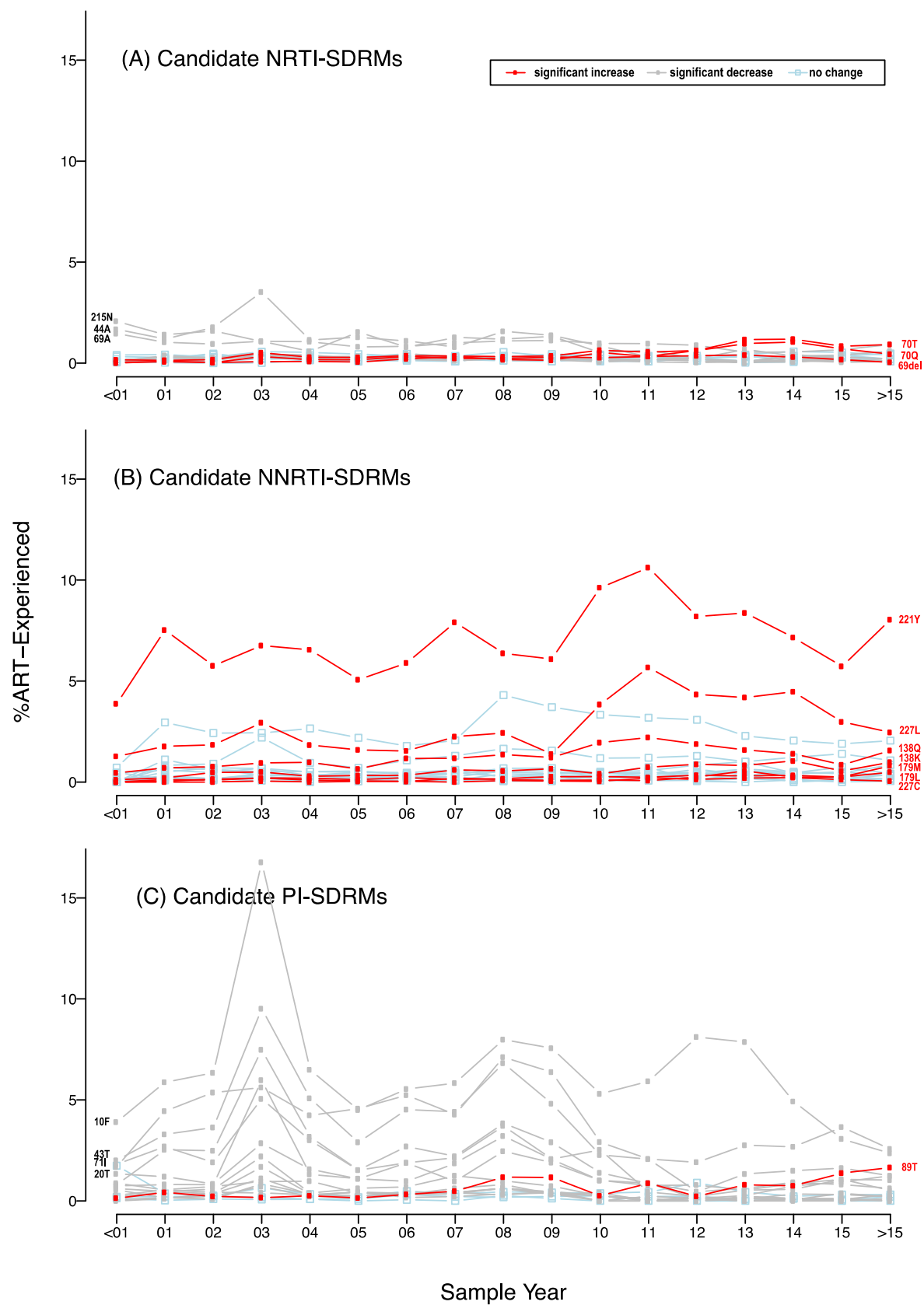


Figure S2. Yearly prevalence of (A) candidate NRTI-SDRMs in NRTI-experienced persons, (B) candidate NNRTI-SDRMs in NNRTI-experienced persons and (C) candidate PI-SDRMs in PI-experienced persons. Candidate SDRMs significantly increased in prevalence are shown in red and indicated in red font on the right of the plotted lines. Candidate SDRMs significantly decreased in prevalence are shown in grey and indicated in black font on the left of the plotted lines. Candidate SDRMs did not change significantly in prevalence are shown in light blue. For candidate SDRMs that decreased with a very low overall prevalence or did not change in prevalence, the labels are not shown. Full lists of candidate NRTI-SDRMs, NNRTI-SDRMs and PI-SDRMs according to the change in prevalence are shown in Table 2, 4, and 6, respectively.