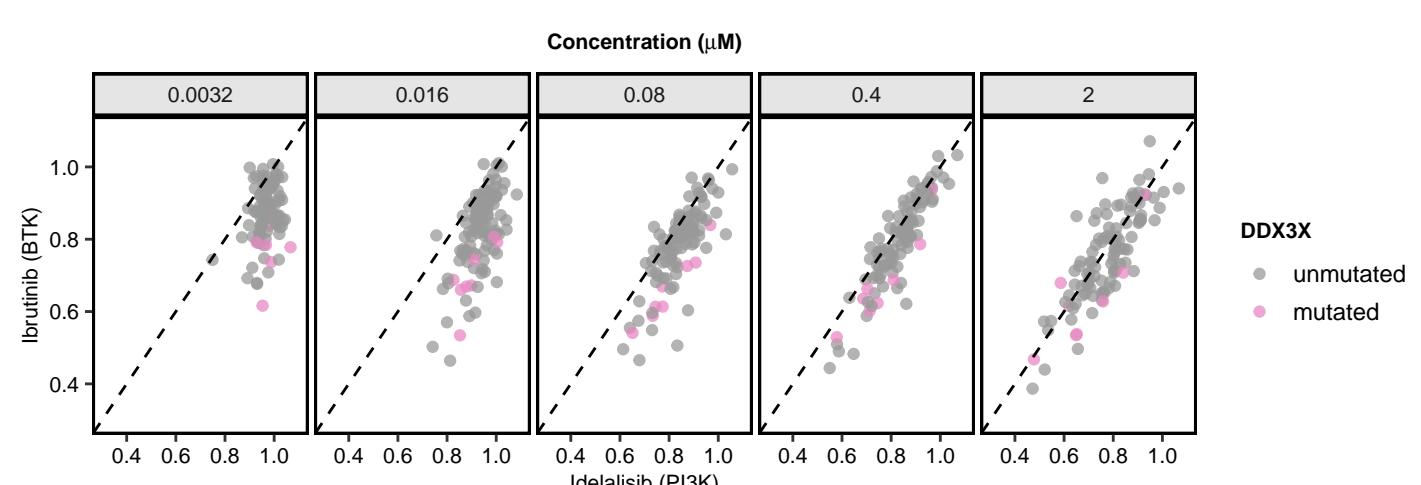


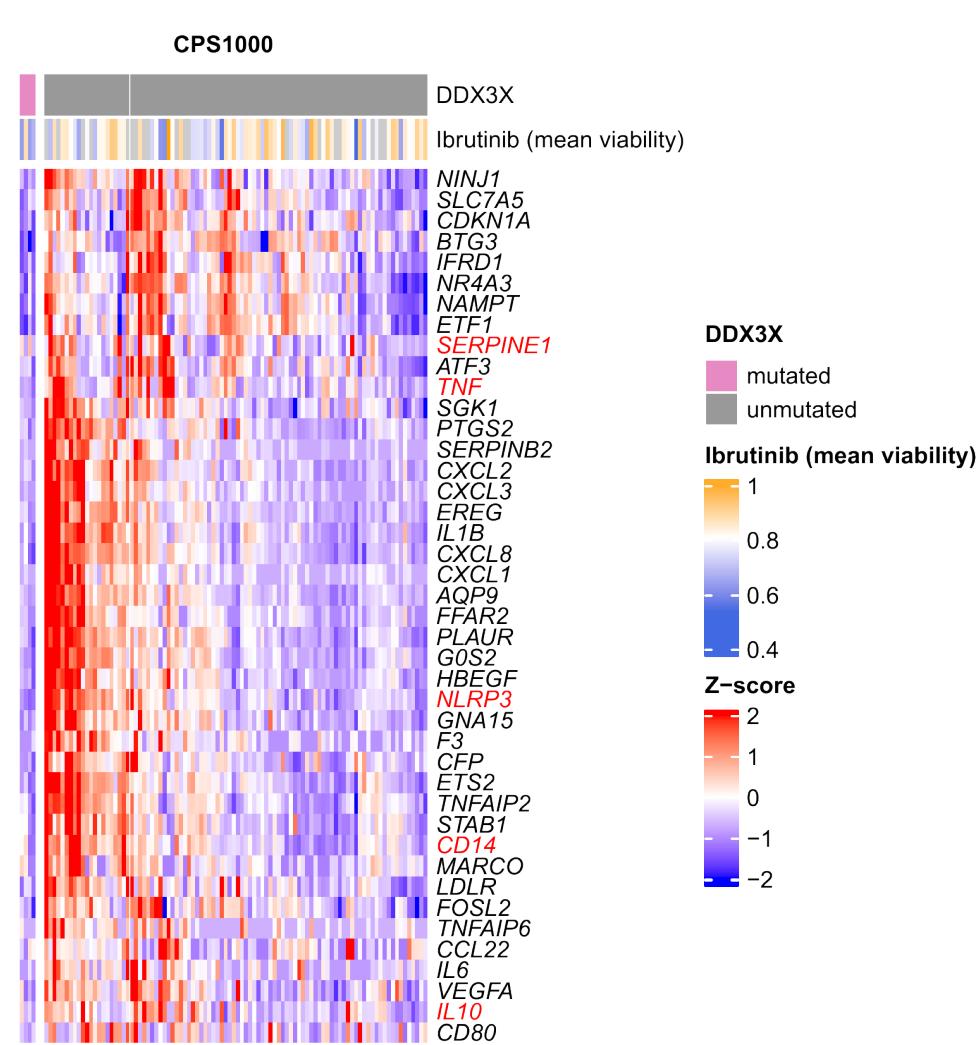
A

ID	Sex	IGHV	DNA mutation	Amino acid alteration	Exon	Type of alteration	Allele frequency
P0293	M	U	c.845dupG	p.C282fs	9	Frameshift insertion	0.86
P0538	M	U	c.1115_1122del	p.E372fs	10	Frameshift deletion	1.00
P0696	F	U	c.A991G	p.M331V	10	Nonsynonymous SNV	0.33
P0711	M	U	c.C1229G	p.S410C	12	Nonsynonymous SNV	0.17
P0880	F	U	c.C1184T	p.T395I	11	Nonsynonymous SNV	0.18
P0987	M	U	c.C1120T	p.Q374X	10	Stopgain	0.45
P1027	M	U	c.T1256G	p.L419R	11	Nonsynonymous SNV	0.94
P1075	M	U	c.C625T	p.Q209X	6	Stopgain	0.96

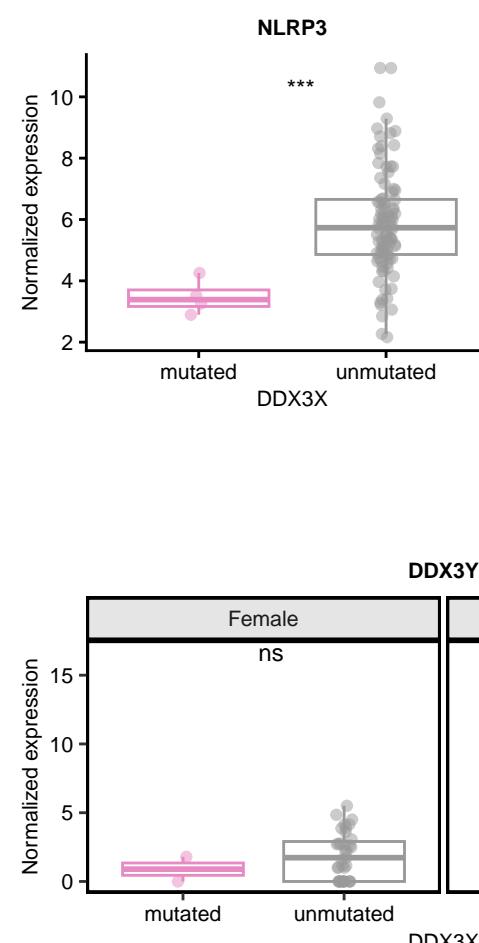
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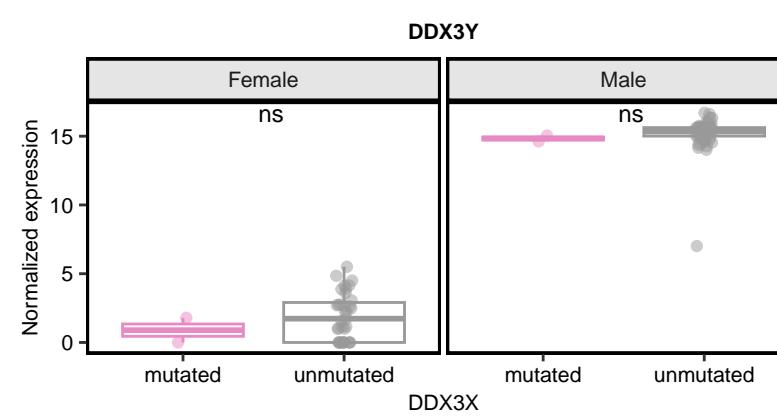
C



E



F



D

