Supplementary Table S2. Discovery phage display screening results for isolating SEP-specific scFvs.

Two naïve phage display Discovery libraries (AXL40 and AXL41) were independently screened against five SEP-containing peptides spanning the exon 2/4 splice junction of CENP-A- Δ Exon3, as per Fig. 1B. 88 single clones from each of the 10 screens were expressed as secretory scFvs and tested in cell supernatant ELISAs against the corresponding SEP and NAT peptides as well as NeutrAvidin alone. "# of Hits" was determined by OD450 \geq 0.2 against the peptide with \geq 2-fold signal over NeutrAvidin. "# of SEP-Specific Hits" was determined by three criteria: (1) OD450 \geq 0.2 against the SEP peptide with \geq 2-fold signal over NeutrAvidin, (2) < 5-fold signal against the NAT peptide over NeutrAvidin, and (3) \geq 2-fold signal against the SEP peptide over the NAT peptide. A subset of SEP-specific hits was sequenced and unique clones were identified both within and between screens. A subset of unique scFvs were purified and tested in antigen and/or scFv titration ELISAs against the corresponding SEP and NAT peptides. Clones with \geq 5-fold background-corrected signal against the SEP peptide over the NAT peptide for at least two titration points were classified as SEP-specific (two examples are shown in Fig. 2A-B, left).

Discovery Screening Results

Discovery Screen # Post		Discovery Screening Results										
P3494		Discovery	ELISA Antigen	# of Hits	# of SEP- Specific	Hits that are	Specific Hits	SEP-Specific Hits (within		in Other Screens	scFv Protein	in Titration
NAT1 57	AXL40	p3494	SEP1	74	- 21	28%	4	3	75%	0	3	3
P3495			NAT1	57								
NAT1 59 59 70 70 70 70 70 70 70 7		p3495	SEP2	74	15	20%	1	1	100%	0	1	1
P3496 SEP3 67 25 37% 20 13 65% 3 [p3497 & p3498 (x2)] 9 2			NAT1	59								
NAT2 48		p3496 -	SEP3	67	- 25	37%	20	13	65%		9	2
P3497 NAT2 59 24 30% 22 6 27% [p3496 & p3498 (x2)] 4 1 P3498 SEP5 57 NAT2 30 25 44% 19 6 32% 3 [p3496 & p3497 (x2), p3496 (x1)] 3 1 P3500 SEP1 0 0 N/A N			NAT2	48								
NAT2 59		p3497	SEP4	81	24	30%	22	6	27%		4	1
P3498			NAT2	59								
P3500 SEP1 0 0 N/A N		p3498	SEP5	57	25	44%	19	6	32%		3	1
P3500 NAT1 2 0 N/A			NAT2	30								
P3500 NAT1 2 0 N/A												
NAT1 2	AXL41	p3500	SEP1	0	0	N/A	N/A	N/A	N/A	N/A	N/A	N/A
P3501 NAT1 0 0 N/A			NAT1	2								
NAT1 0 SEP3 71 64 90% 57 32 56% 1 [p3504 (x1)] 16 2 2 2 3 3 3 44 96% 27 11 41% 0 1 0 1 0 1 0 2 2 2 2 55% 4 3 75% 1 [p3502 (x1)] 0 N/A		p3501	SEP2	0	0	N/A	N/A	N/A	N/A	N/A	N/A	N/A
P3502 NAT2 10 64 90% 57 32 56% 1 [p3504 (x1)] 16 2 p3503 SEP4 46 A4 96% 27 11 41% 0 1 0 p3504 SEP5 22 12 55% 4 3 75% 1 [p3502 (x1)] 0 N/A			NAT1	0								
P3503 SEP4 46 44 96% 27 11 41% 0 1 0 0 0 0 0 0 0 0		p3502	SEP3	71	64	90%	57	32	56%	1 [p3504 (x1)]	16	2
p3503 NAT2 3 44 96% 27 11 41% 0 1 0 0 1 0 p3504 SEP5 22 12 55% 4 3 75% 1 [p3502 (x1)] 0 N/A			NAT2	10								
NAT2 3		p3503 -	SEP4	46	44	96%	27	11	41%	0	1	0
p3504 12 55% 4 3 75% 1 p3502 (x1) 0 N/A			NAT2	3								
NAT2 12 12 17 18 18 18 18 18 18 18 18 18 18 18 18 18		p3504 -	SEP5	22	12	55%	4	3	75%	1 [p3502 (x1)]	0	N/A
10.112			NAT2	12								