

BD[®] OMICS-One Protein Panels

The power of RNA + protein without the high cost and complexity



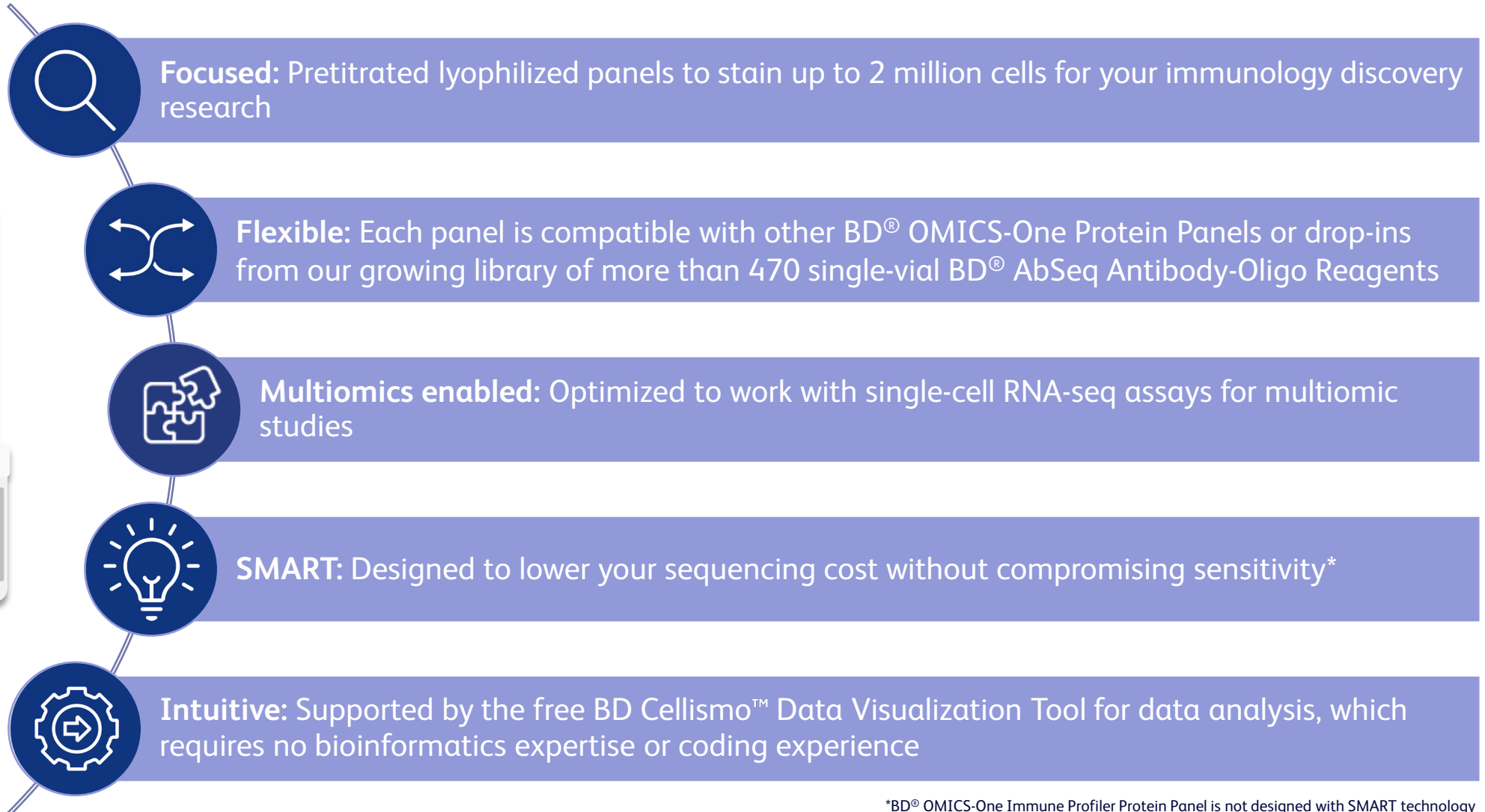
BD[®] OMICS-One Protein Panels for CITE-seq

CITE-seq helps to unlock deeper insights about your samples and can reveal novel cell types and different cell states.

We offer a complete CITE-seq solution with a streamlined sample-to-data workflow. Now with predesigned BD[®] OMICS-One Protein panels, CITE-seq on the BD Rhapsody[™] System is even more accessible.



BD® OMICS-One Protein Panels

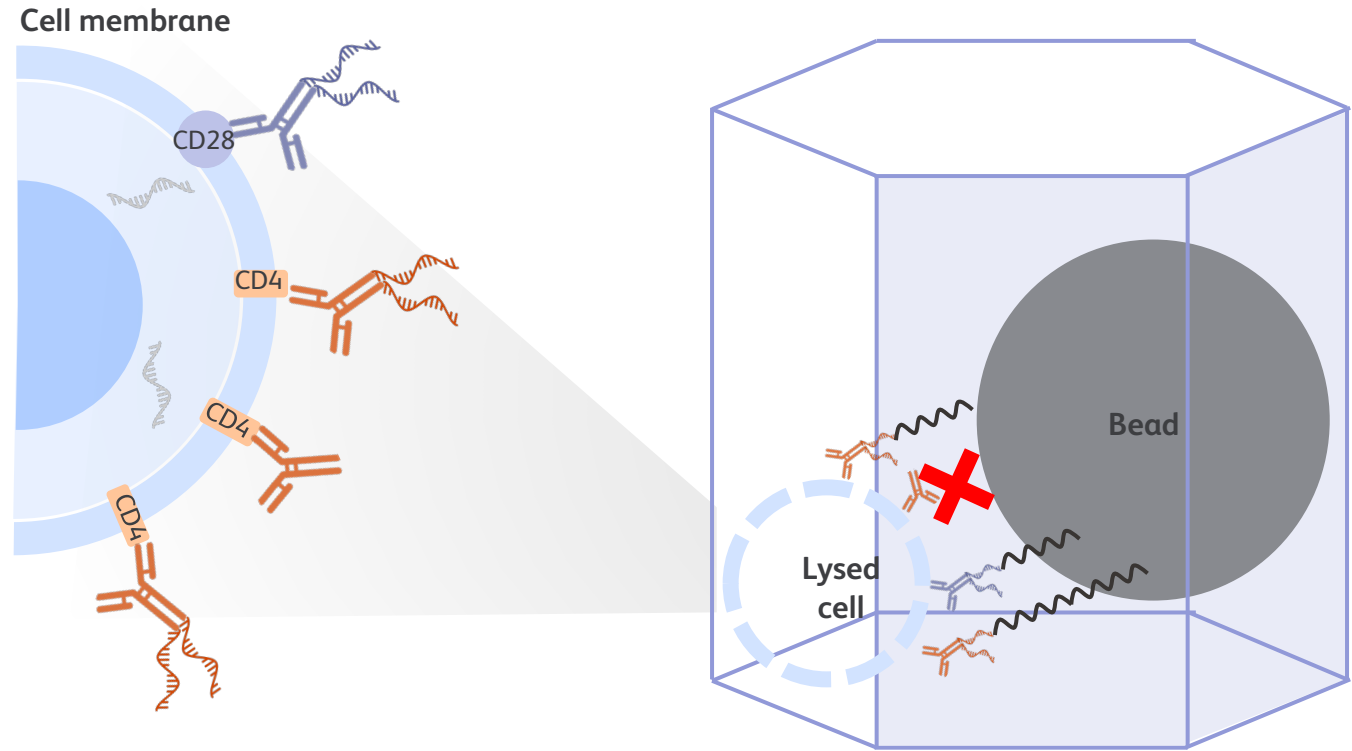


*BD® OMICS-One Immune Profiler Protein Panel is not designed with SMART technology

Manage sequencing costs and improve sensitivity with SMART panel design

SMART panel design helps lower sequencing costs while increasing data resolution utilizing carefully selected, pretitrated concentrations of antibody-oligos (SMART-titrated antibodies) against abundant primary markers and allowing re-allocation of sequencing reads to markers expressed at lower levels.

With SMART panel design, now low-expression markers can be quantified without having to do deeper sequencing and incurring high sequencing costs.



More NGS sequencing reads  allocated to low expressors

Regular CITE-seq panel

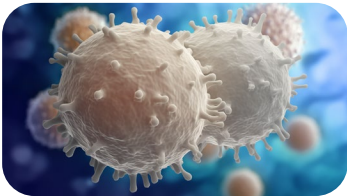


With SMART panel design

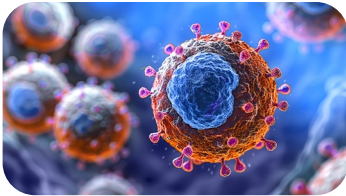


Available panels

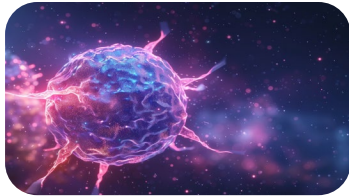
Modular panels



[T-Cell Protein Panel](#)



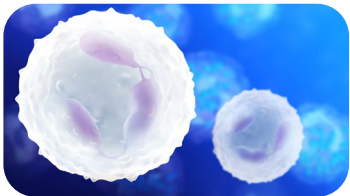
[B-Cell Protein Panel](#)



[Tumor Protein Panel](#)

NEW

Focused integration panels



[Adaptive Protein Panel](#)

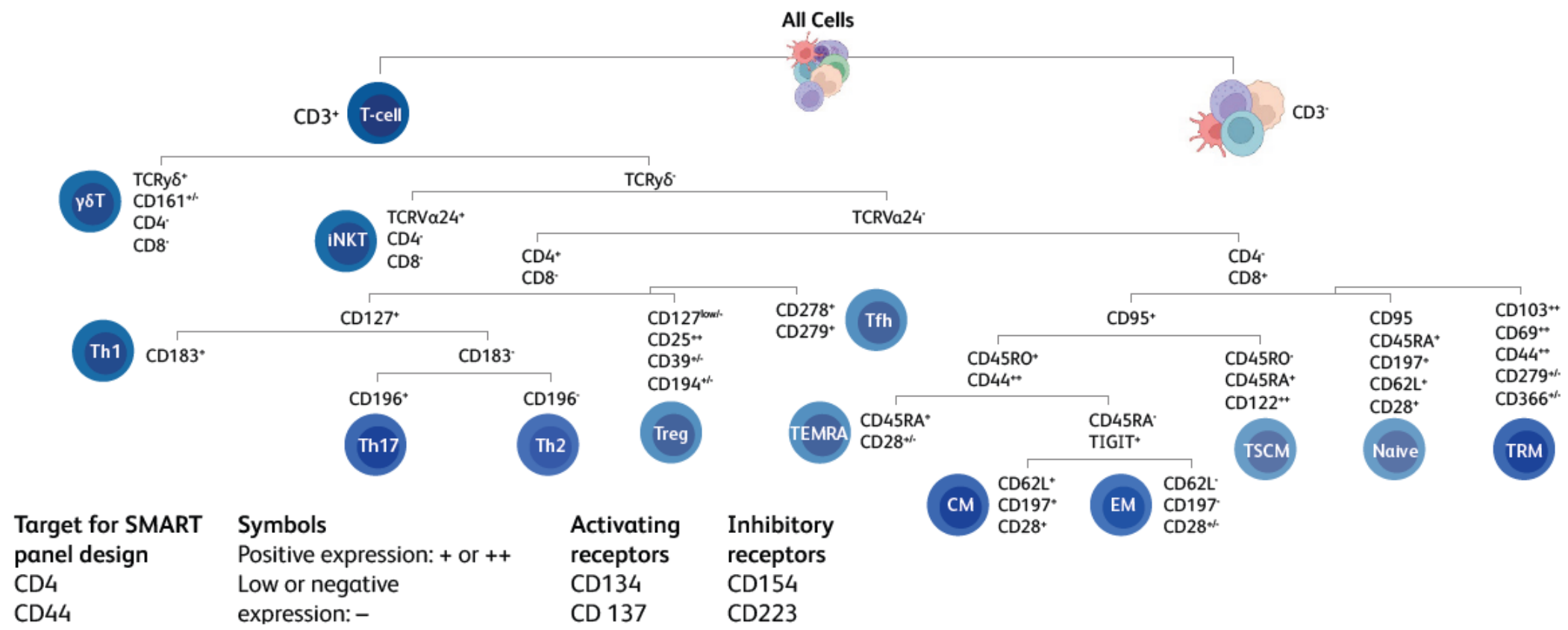


[Immuno-Oncology Protein Panel](#)

NEW

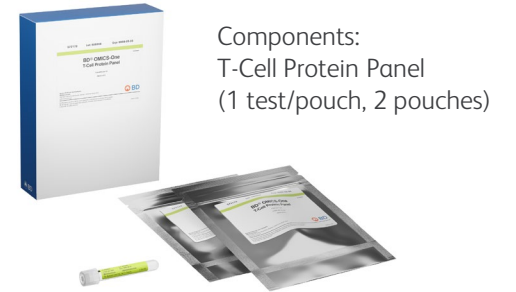
BD[®] OMICS-One T-Cell Protein Panel

Explore T-cell populations and their activation and suppressed states



T-cell populations and activation states monitored by this panel.

BD® OMICS-One T-Cell Protein Panel specificities



Specificity	Clone
CD3	UCHT1
CD4*	SK3
CD8	SK1
CD25	2A3
CD28	L293
CD44*	L178
CD45RO	UCHL1
CD45RA	HI100
CD69	FN50
CD62L	DREG-56

Specificity	Clone
CD95	DX2
CD103	BER-ACT8
CD127	HIL-7R-M21
CD134	ACT35
CD137	4B4-1
CD154	TRAP1
CD161	HP-3G10
CD183	1C6/CXCR3
CD194	1G1
CD196	11A9

Specificity	Clone
CD197	2-L1-A
CD223	T47-530
CD272	J168-540
CD278	DX29
CD279	EH12.1
CD357	V27-580
CD366	7D3
TCR γ / δ	11F2
TCR V α 24-J α 18	6B11
TIGIT	TgMab-2

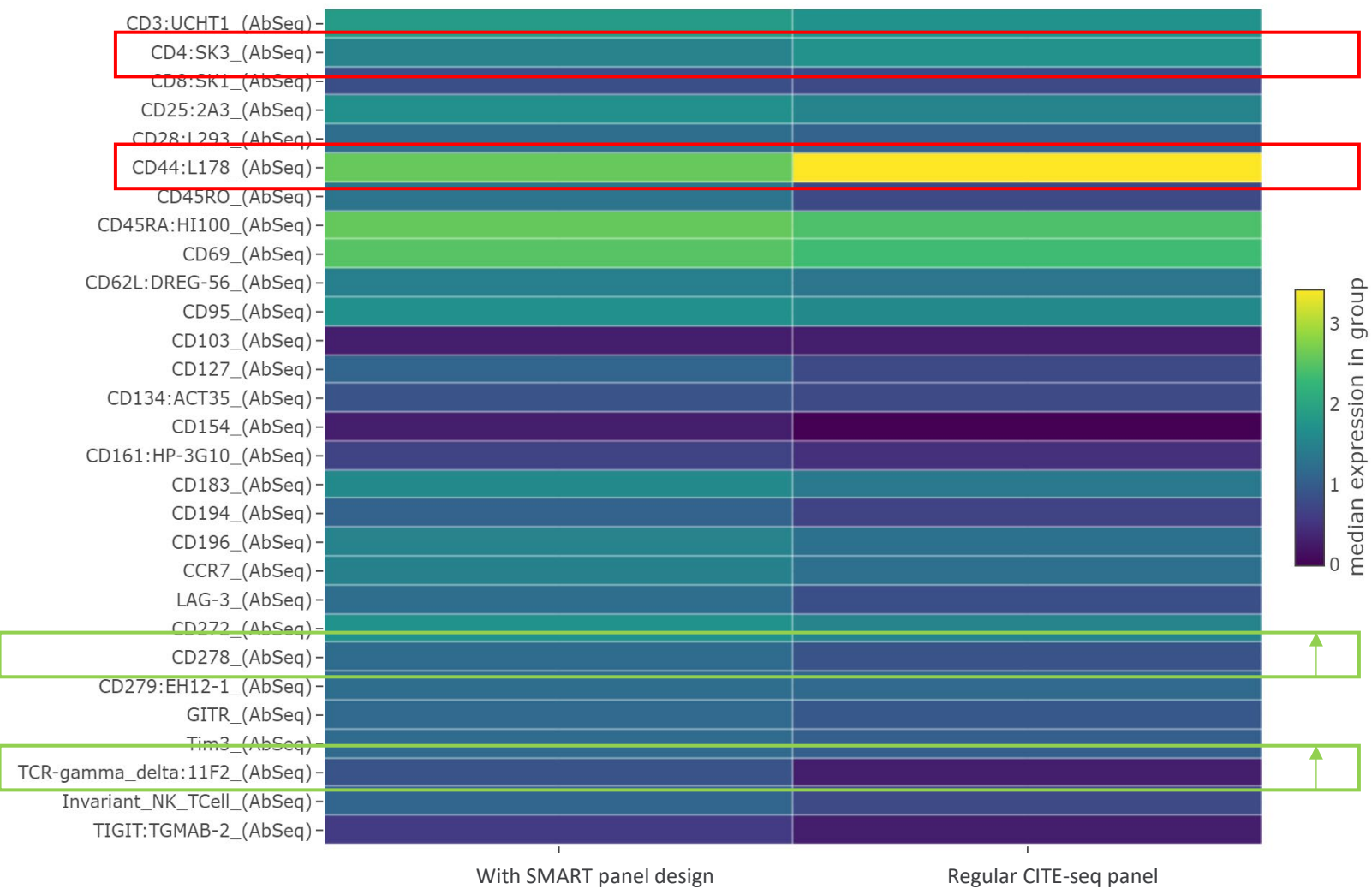
*Targets for SMART panel design

Fewer sequencing reads consumed by CD4 and CD44 and more reads allocated to lowly expressed markers

Percent of Total Sequencing Reads Consumed		
Markers	Without SMART panel design	With SMART panel design
Reduction of sequencing reads allocated to primary markers ▼		
CD4	5.92	4.07
CD44	46.10	8.70
Read re-allocation to lowly expressed markers ▲		
CD3	2.30	4.25
CD8	0.98	1.59
CD25	3.72	7.52
CD45RO	0.87	1.48
CD45RA	3.88	9.17
CD69	9.50	19.07
CD62L	1.35	2.61
CD95	1.18	1.86
CD127	0.61	1.10
CD137	1.64	3.14
CD197	0.91	1.76
CD223	0.78	1.71

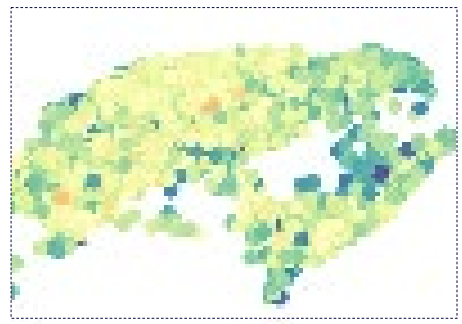
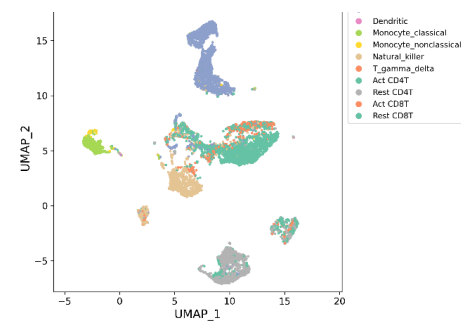
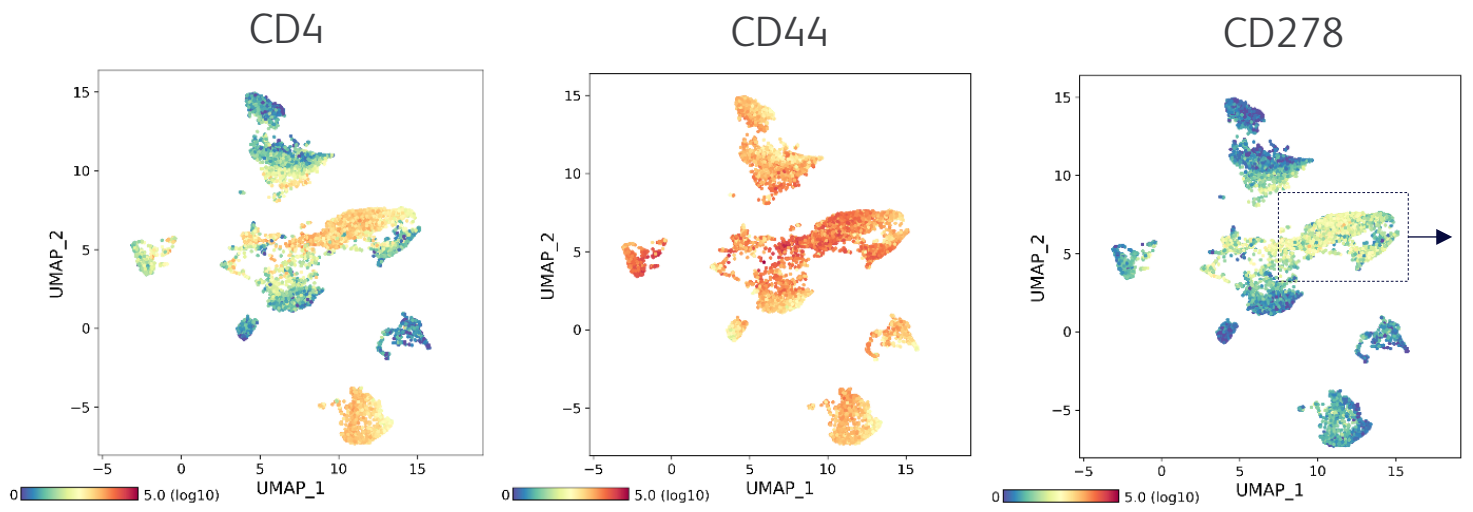
Percent of Total Sequencing Reads Consumed		
Markers	Without SMART panel design	With SMART panel design
CD279	1.77	2.67
CD366	1.80	3.39
CD103	0.23	0.46
CD134	0.88	1.28
CD154	0.15	0.29
CD161	0.38	0.64
CD183	4.87	4.15
CD194	1.94	3.79
CD196	2.65	4.16
CD272	1.72	3.34
CD278	0.65	1.33
CD28	0.95	1.68
CD357	0.57	1.23
TCR Vα24-Jα18	0.86	1.93
TCRγ/δ	0.34	0.79
TIGIT	0.38	0.74

Increased AbSeq molecule detection of low expressors

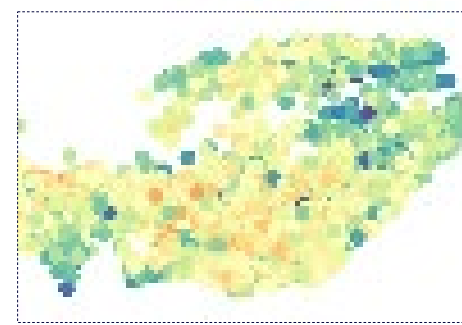
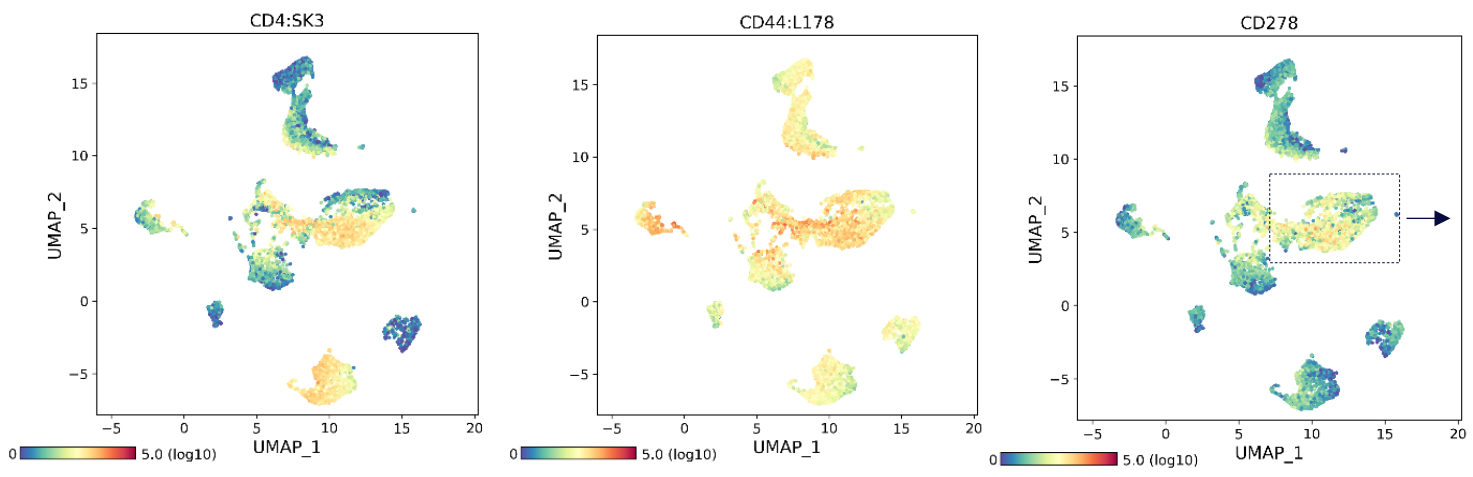


CD4 and CD44 detection is not compromised, while better resolution of low expressors is found with SMART panel design

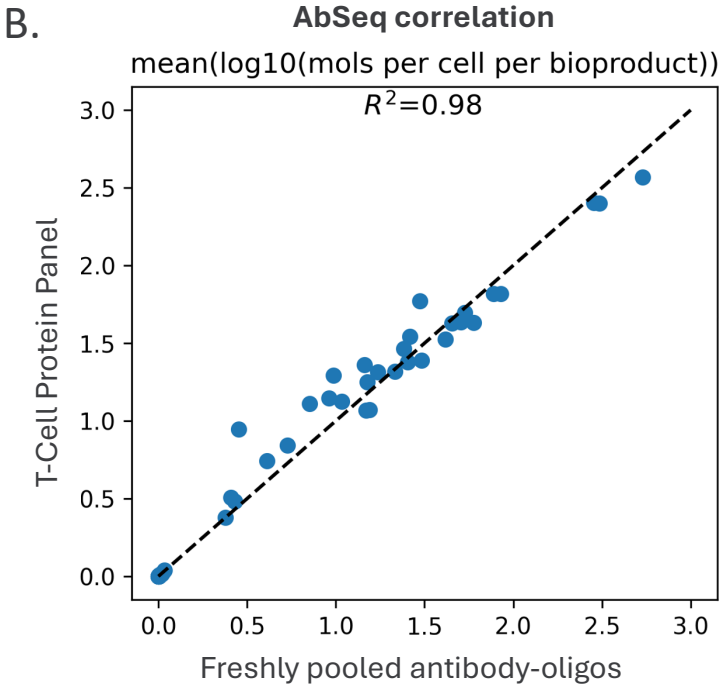
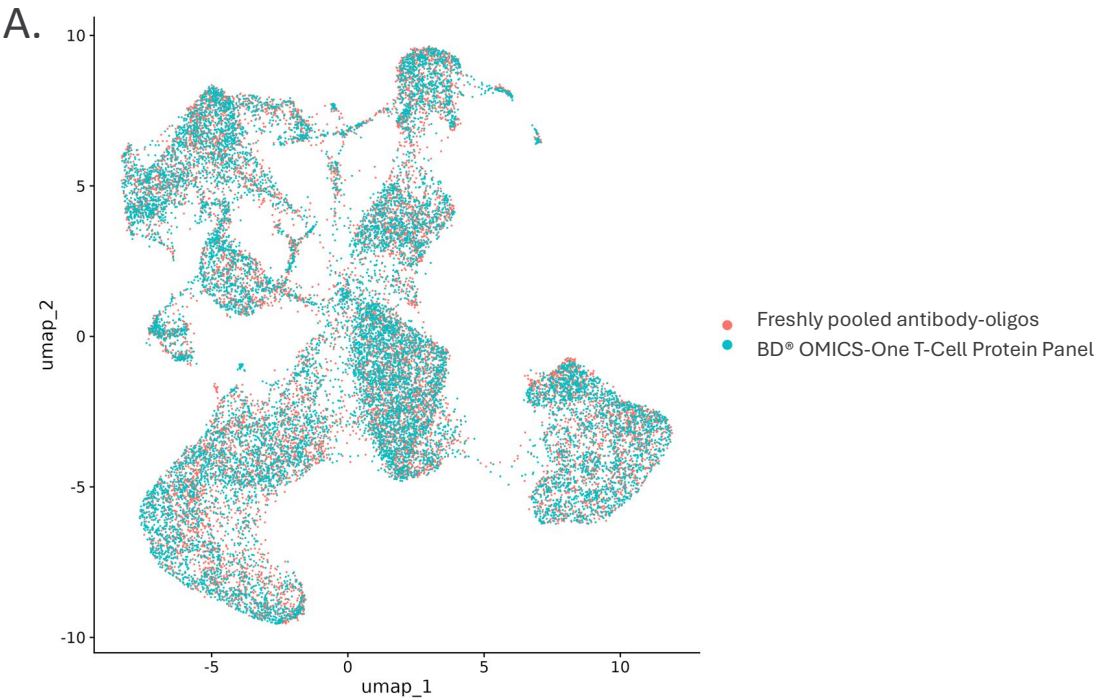
Without SMART panel design



With SMART panel design

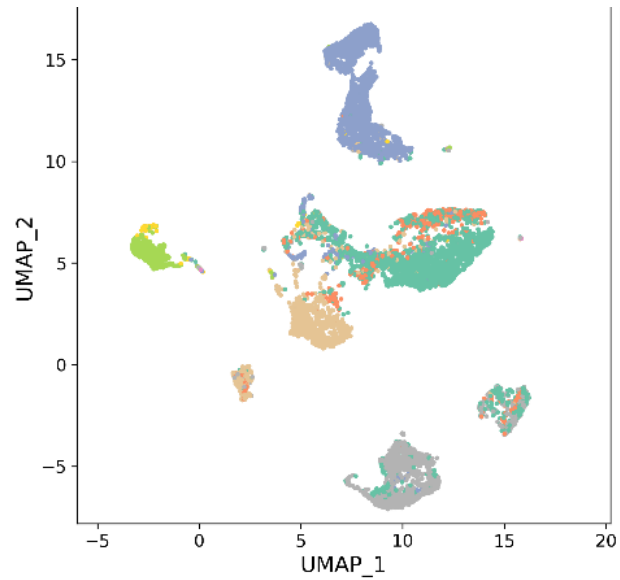


Comparable performance of the lyophilized panel to freshly pooled BD[®] AbSeq Antibody-Oligo Reagents

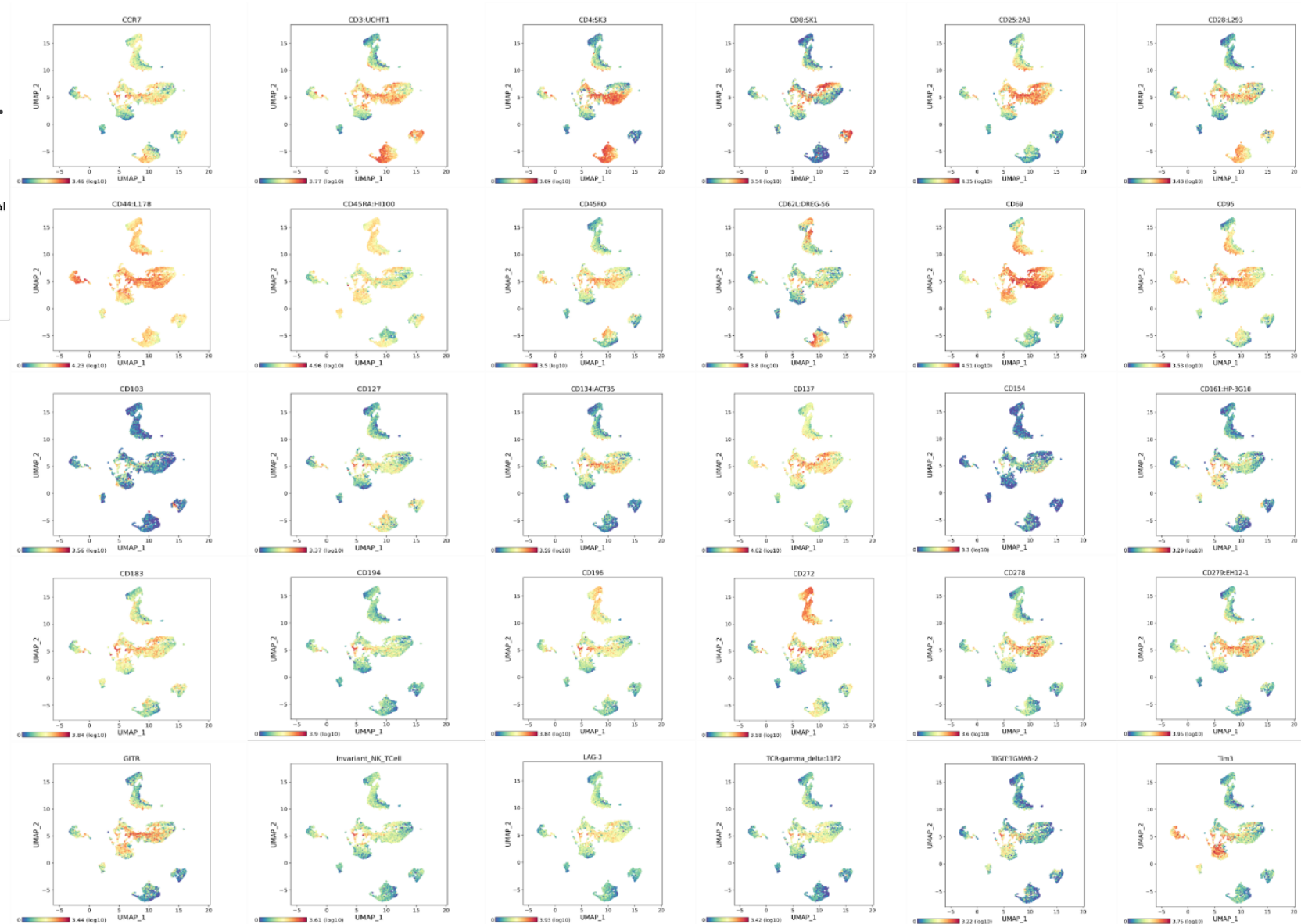


Detect 30 critical T-cell markers in your samples with confidence

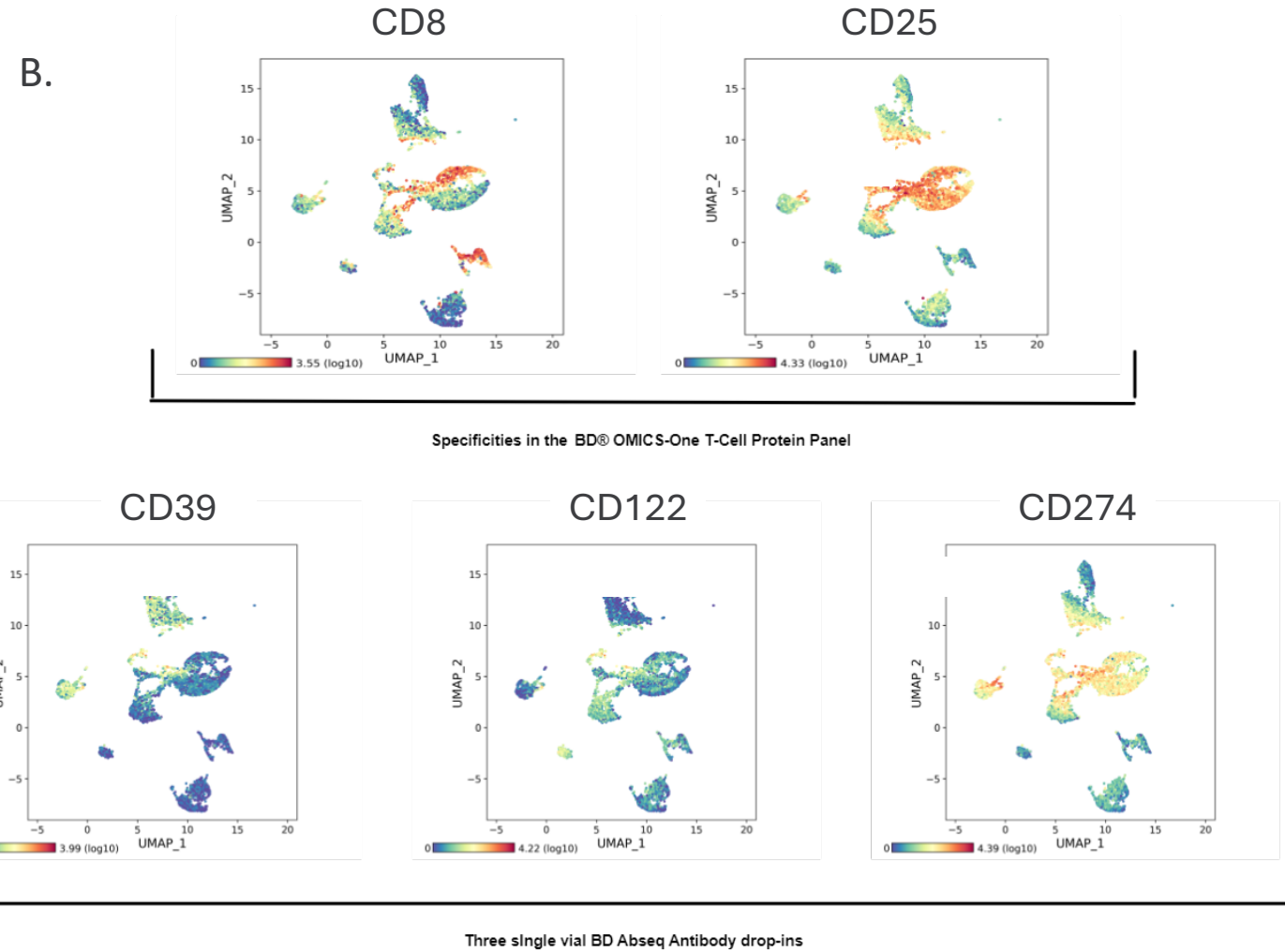
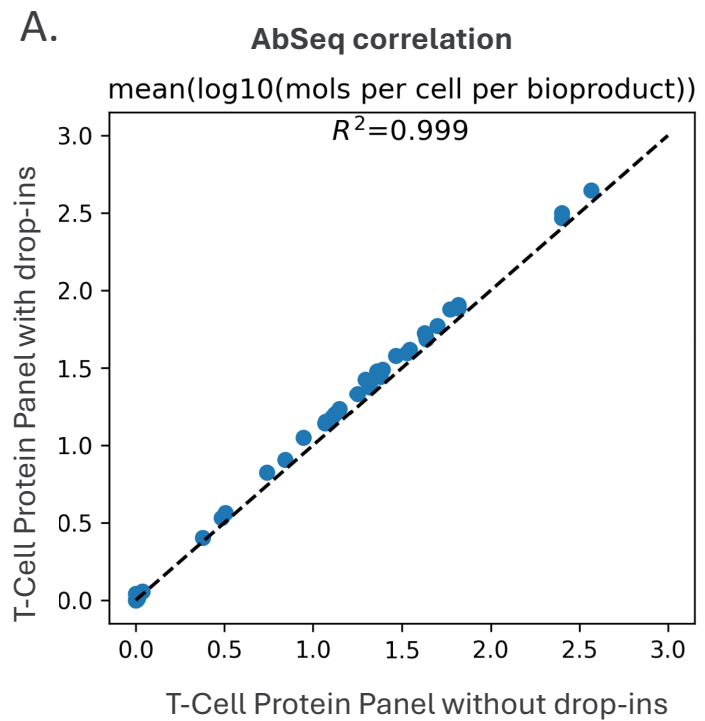
A.



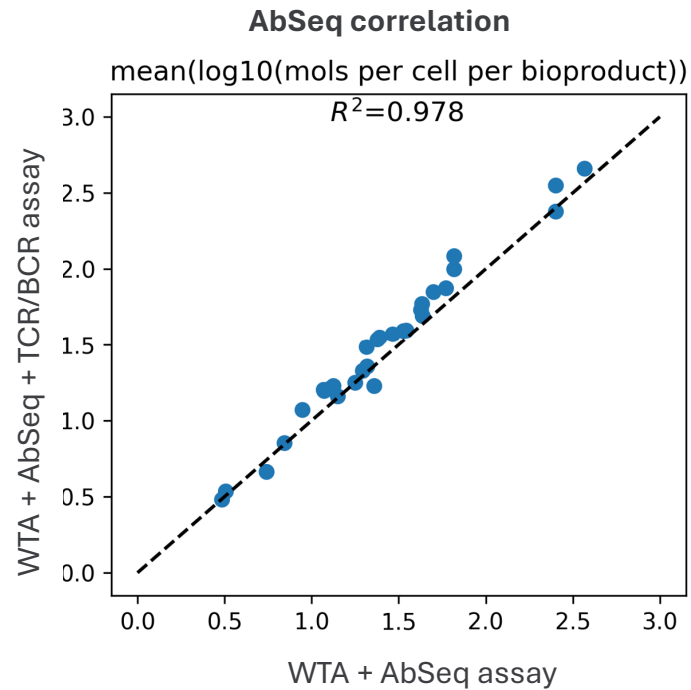
B.



Flexible: Add additional specificities of interest to the BD® OMICS-One T-Cell Protein Panel without compromising performance

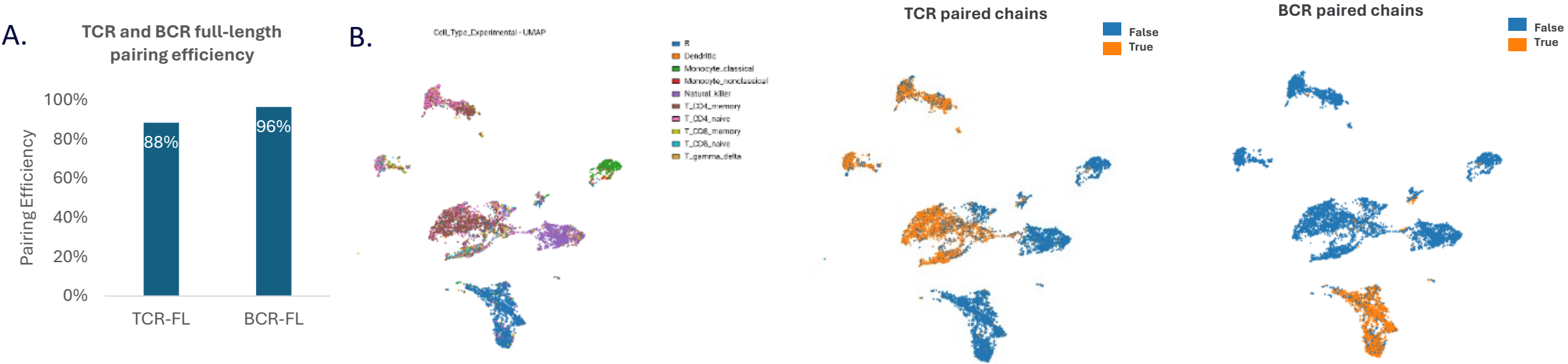


Multiomics enabled: The BD[®] OMICS-One T-Cell Protein Panel is designed to work with whole transcriptome and TCR/BCR profiling assays



The addition of the BD Rhapsody[™] TCR/BCR Next Assay does not impact BD[®] OMICS-One T-Cell Protein Panel performance

Multiomics enabled: The BD[®] OMICS-One T-Cell Protein Panel is designed to work with whole transcriptome and TCR/BCR assays

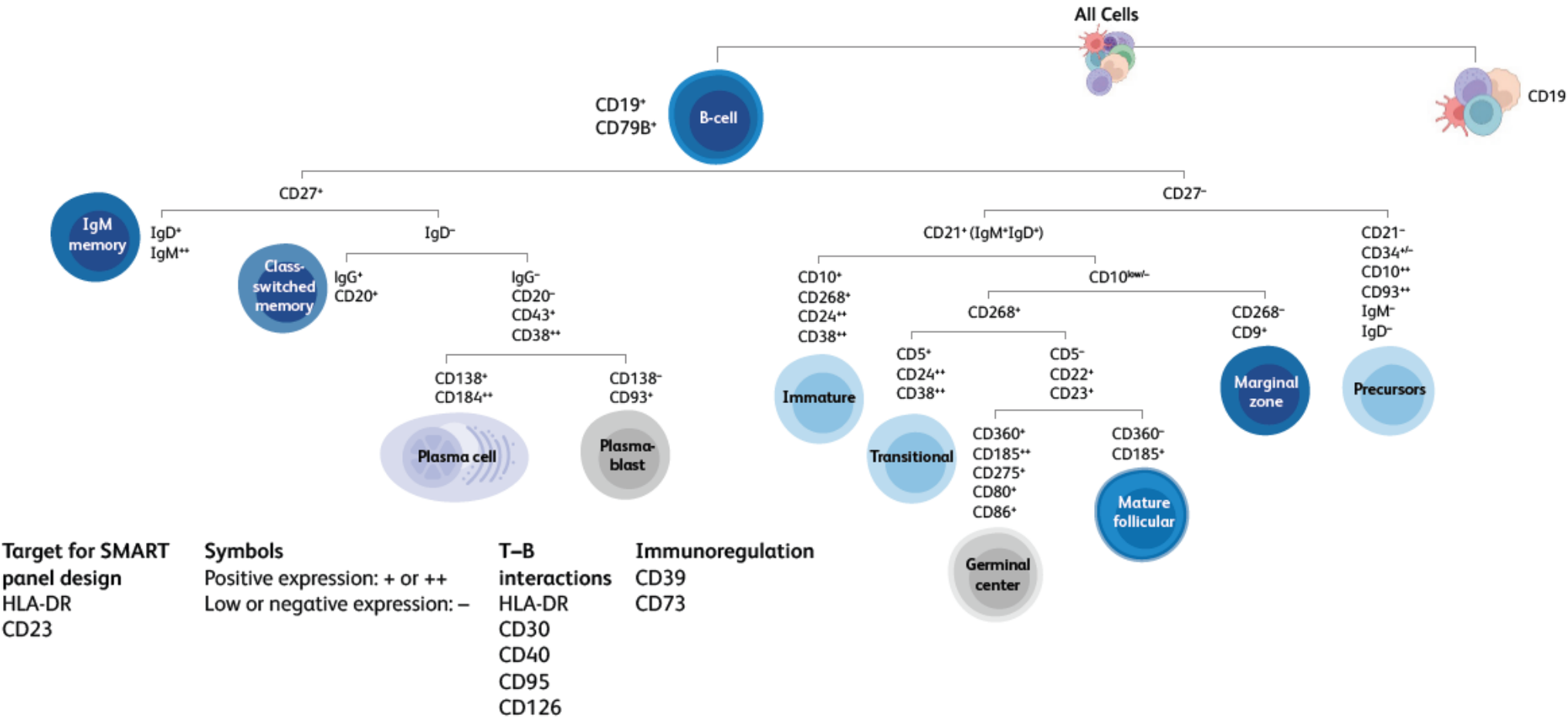


C. ● FWR1 ● CDR1 ● FWR2 ● CDR2 ● FWR3 ● CDR3 ● FWR4

Cell Index	TCR Chain	VDJ Translation Trimmed
1057	TCR_Alpha	AQTVTQSQPEMSVQEAETVLSCTYD TS ENNYLFWYKQPPSRQ MIL VIR Q EAYK Q Q NA TENRFSVNFQKA AK SFSLKISDSQLGDTAM Y FCALPGSE NN DM R FGAG TR LTVKP
1057	TCR_Beta	NAGVTQTPKFRILKIGQSM T LQCTQD MN HNMYWYRQDPGMGLKLIY S VGAGITDKGEVPNGYNVSRST ED FPLRLELAAPSQTSVYFCASSV V GRGY FG SG TR LTVV
6784	TCR_Alpha	GQ N IDQPT EM TATEGAIVQINCTYQ TS GF N GLFWYQQHAGEAPT FL S N VL D GLEEKGRFSSFLSRSGYSYLL K ELQMKDSASYLC AV KGSYIPT FG RGTSLIVHP
6784	TCR_Beta	DAGVIQSPRHEVTEMGQEVTLRCKPIS GH NSLFWYRQTMMRGLELLI Y FNN N VPIDDSGMPEDRFS AK MPNASFSTLKI Q PSEPRDSAVYFC AS TFR T GGPETQ Y FGPG TR LLVL
23618	TCR_Delta	AIELVPEHQTPVPSIGVPATLRCSMK GE AIGNYINWYRK T QNTMTFIY R EKDIYGP G KDNFQGDIDIAKNLAVLKILAPSERDEGSY Y CAD T ATGGRSSWD TR QMFFGTG IK LFVEP
23618	TCR_Gamma	AGHLEQPQISSTKTL S K T ARLECV S GISATSVYWYRERPG EV IQFLV S ISYD G TVRKESGIPSGKFEVDRI P ETSTSTLTIHNVEKQDIATDYCALWEVRG Y KKFFG
39145	TCR_Alpha	DAKTQPPSMDCAEGRAANLPCNH S TISGNEVYWYRQ I HSQGPQY I HLK NN ETNEMASLI ED RKSSTLILPHATLRDTAVY C IVRLERGG S NYKLT FG KGTLLTV N P
39145	TCR_Beta	NAGVTQTPKFRVLK T GQSM T LLCAQD MN HEMYWYRQDPGMGLRLI H Y SV EG T TAKGEVPDGY N VSRLKKQ N FL L GLESAAPSQTSVYFC AS RFGSGNTI Y FGEGSWLT V
44437	TCR_Alpha	DQQVKQNSPSLSVQEGRIFYSEL*LYQH V *LPMVQKIP C *RSY P DIK F H* G * R WKIHCFLKQK C QAPLSAHCALPAWRLCSV LL SKQ* H AWSRDQ T DSKT
44437	TCR_Beta	DTGVSQDPRHKITKRGQNVTFRCD P ISEHNRLYWYRQ T LGQGPEFLTY F QNEAQLEKSRLSDRFS A ERP K GSFSTLEIQ R TEQGD S AM Y LCASSVGFTETQ Y FGPG TR LLVL

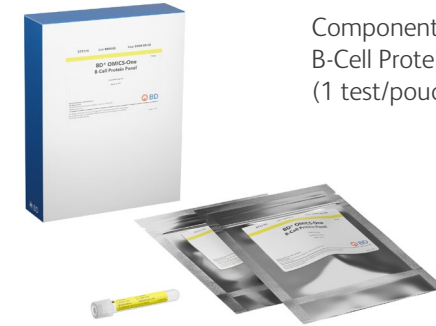
BD[®] OMICS-One B-Cell Protein Panel

Explore B-cell populations and their activation and suppressed states



B-cell populations and activation states monitored by this panel.

BD® OMICS-One B-Cell Protein Panel specificities



Components:
B-Cell Protein Panel
(1 test/pouch, 2 pouches)

Specificity	Clone
CD1d	CD1d42
CD5	UCHT2
CD9	M-L13
CD10	HI10A
CD19	SJ25C1
CD20	2H7
CD21	B-LY4
CD22	HIB22
CD23	EBVCS-5
CD24	ML5

Specificity	Clone
CD27	M-T271
CD30	BERH8
CD34	581
CD38	HB7
CD40	5C3
CD43*	1G10
CD73	AD2
CD79b	CB3-1
CD80	L307.4
CD95	DX2

Specificity	Clone
CD126	M5
CD138	MI15
CD184	12G5
CD185	RF8B2
CD268	11C1
CD275	2D3/B7-H2
HLA-DR*	G46-6
IgD	IA6-2
IgG	G18-145
IgM	G20-127

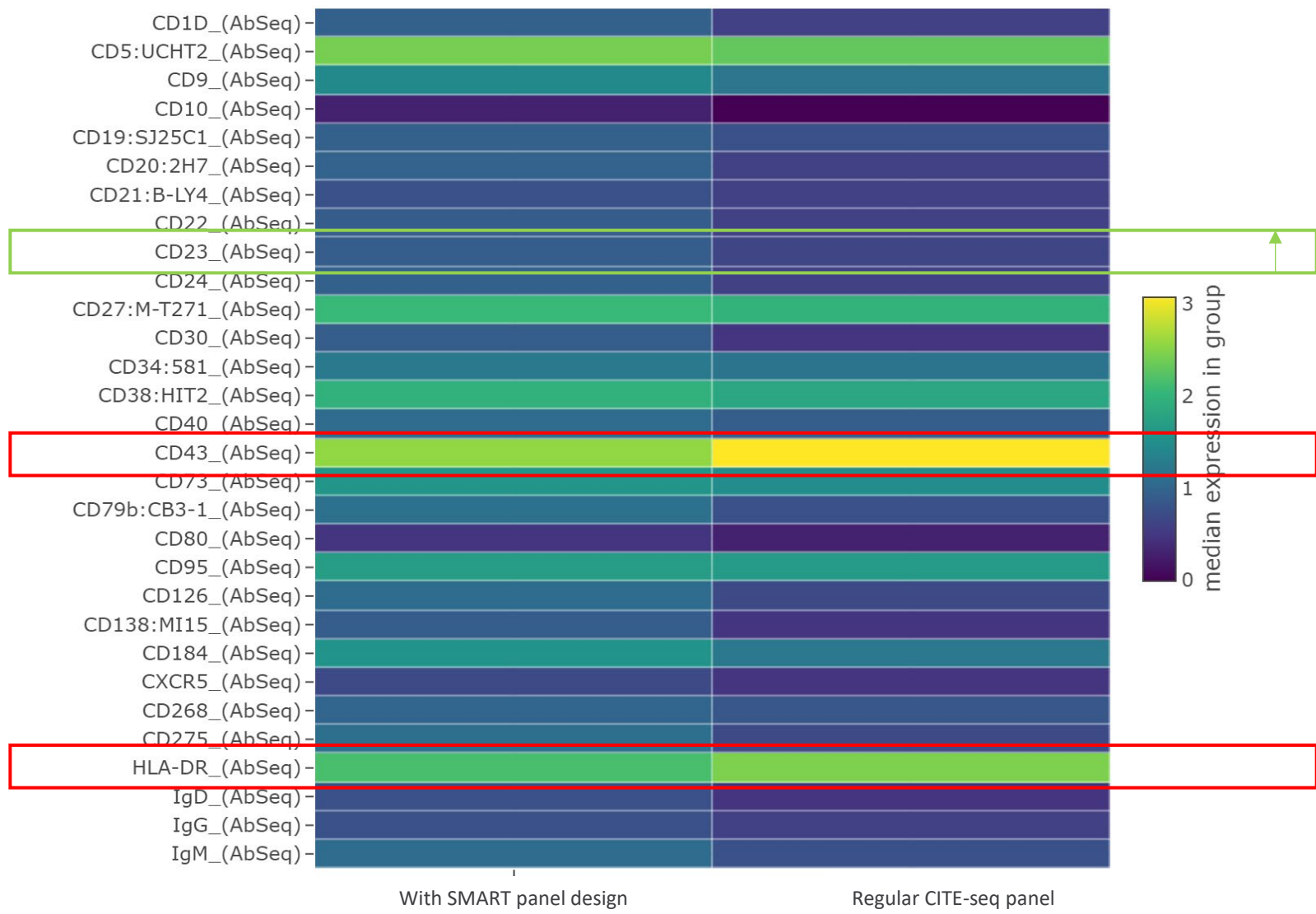
*Targets for SMART panel design

Fewer sequencing reads consumed by CD43 and HLA-DR and more reads allocated to lowly expressed markers

Percent of Total Sequencing Reads Consumed		
Markers	Without SMART panel design	With SMART panel design
Reduction of sequencing reads allocated to primary markers ▼		
CD43	17.09	8.92
HLA-DR	37.76	23.13
Read re-allocation to lowly expressed markers ▲		
CD5	5.74	8.61
CD9	1.76	2.64
CD19	1.22	1.97
CD20	1.28	3.19
CD23	1.72	2.53
CD27	4.05	5.47
CD38	2.14	3.27
CD40	0.93	2.10
CD138	0.41	0.79
CD184	0.88	2.30
IgD	0.87	1.41
IgG	0.66	1.01

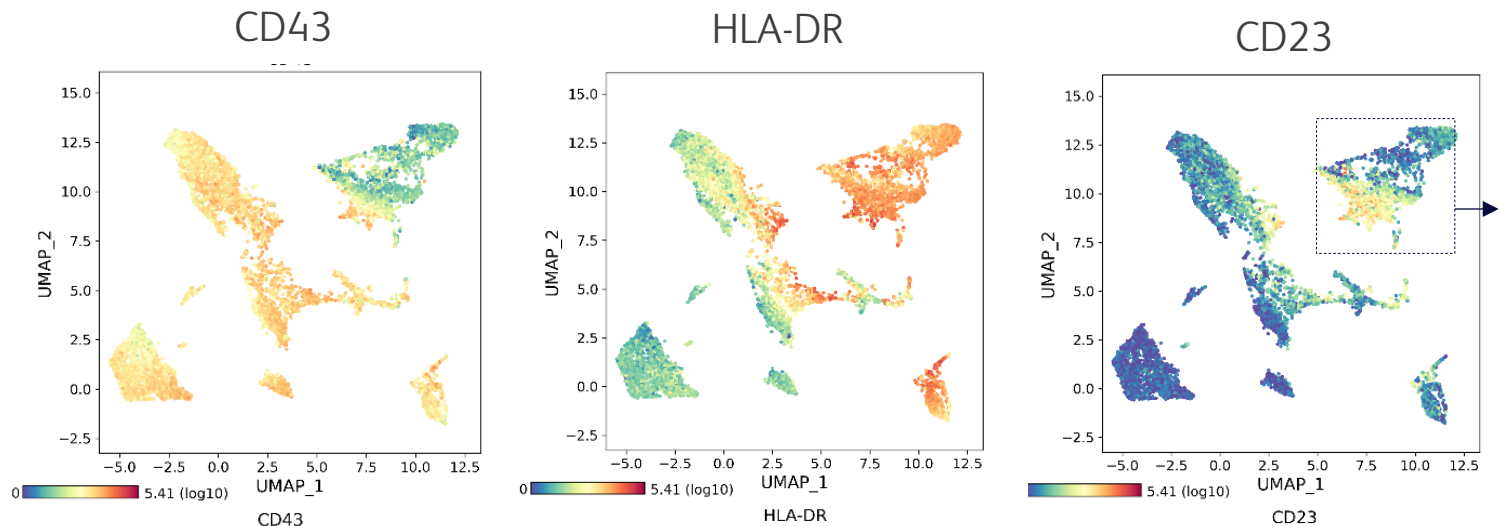
Percent of Total Sequencing Reads Consumed		
Markers	Without SMART panel design	With SMART panel design
IgM	1.71	2.63
CD10	0.21	0.31
CD126	0.51	0.81
CD1d	0.81	1.37
CD21	0.56	0.84
CD22	0.91	1.50
CD24	1.49	2.21
CD268	1.31	1.98
CD275	1.70	3.76
CD30	0.73	1.14
CD34	6.13	5.79
CD73	2.34	3.39
CD79b	2.76	3.85
CD80	0.44	0.60
CD95	1.38	1.66
CD185	0.37	0.72

Increased AbSeq molecule detection of low expressors

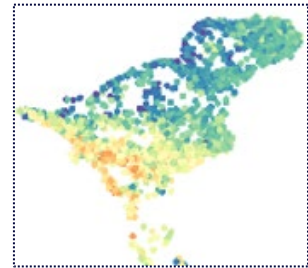
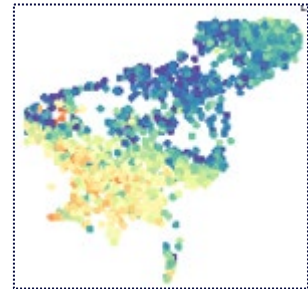
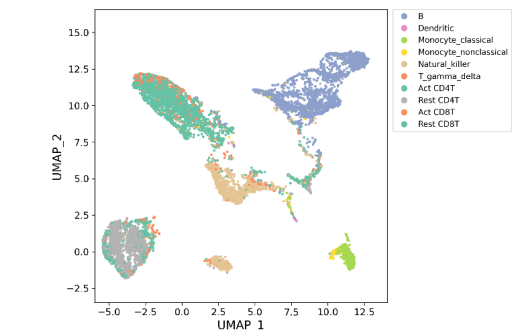
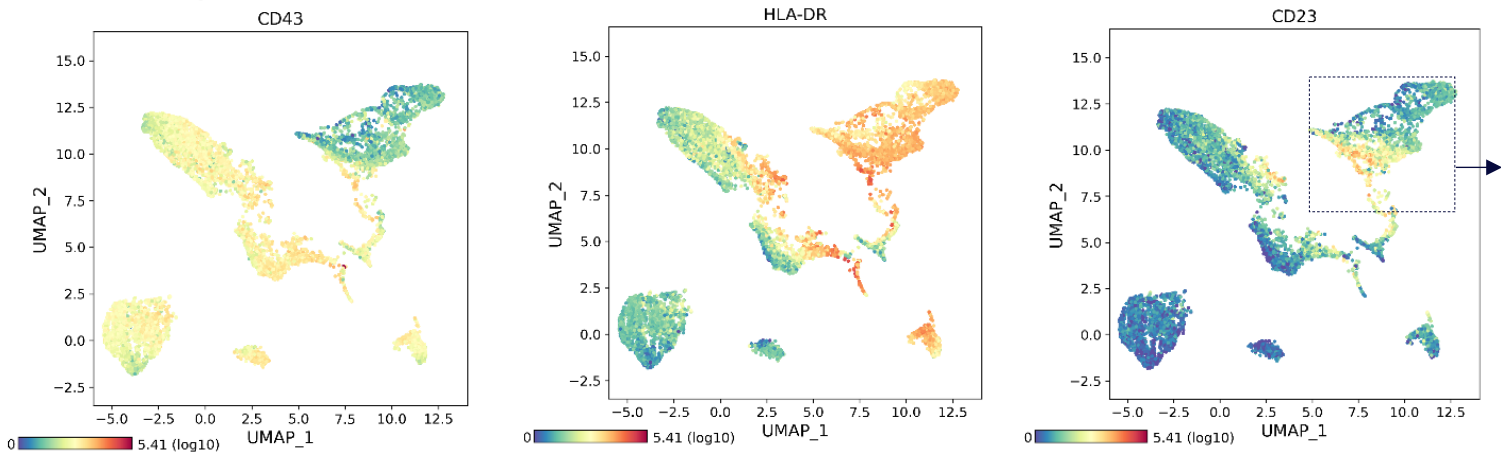


CD43 and HLA-DR detection is not compromised, while better resolution of low expressors is found with SMART panel design

Without SMART panel design

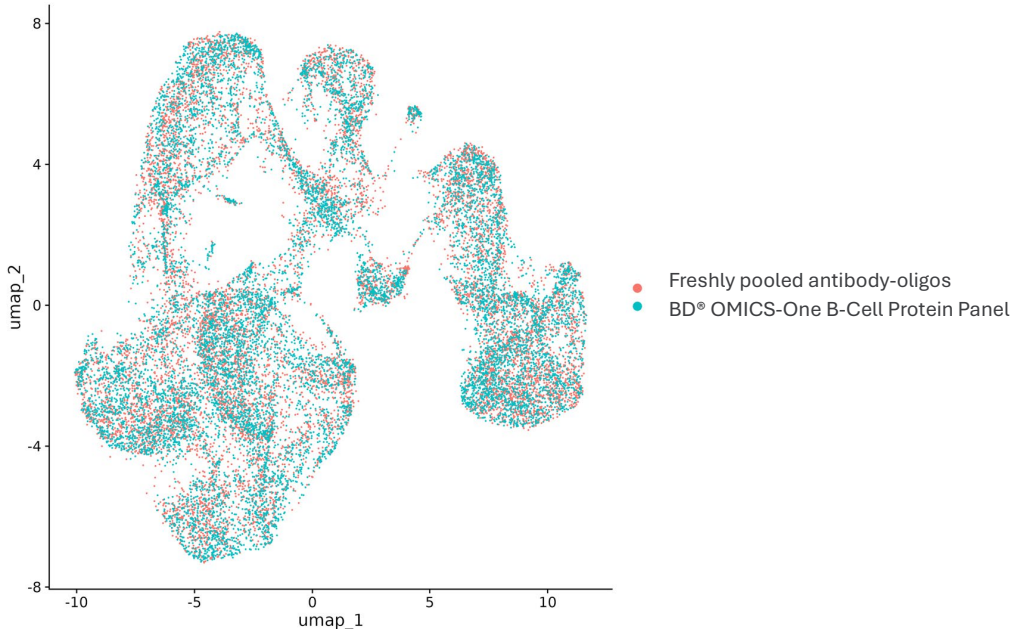


With SMART panel design

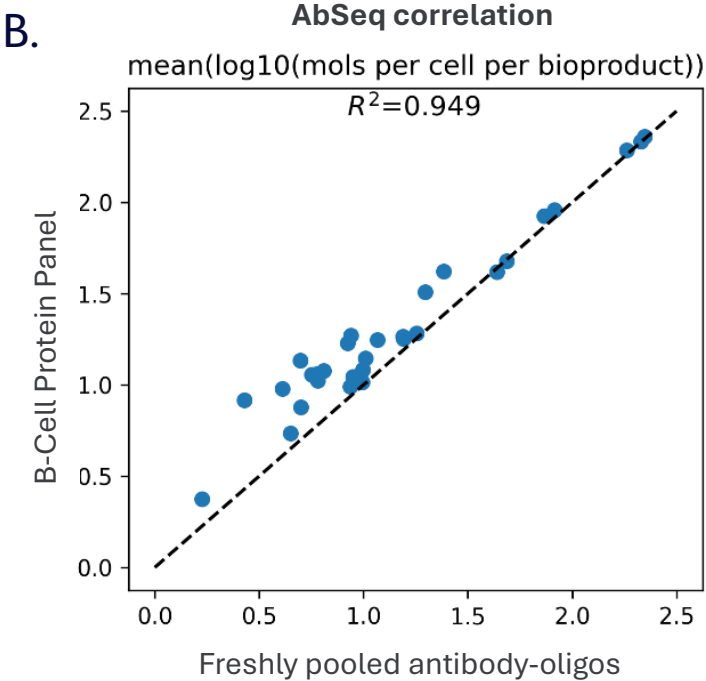


Comparable performance of the lyophilized panel to freshly pooled BD[®] AbSeq Antibody-Oligo Reagents

A.

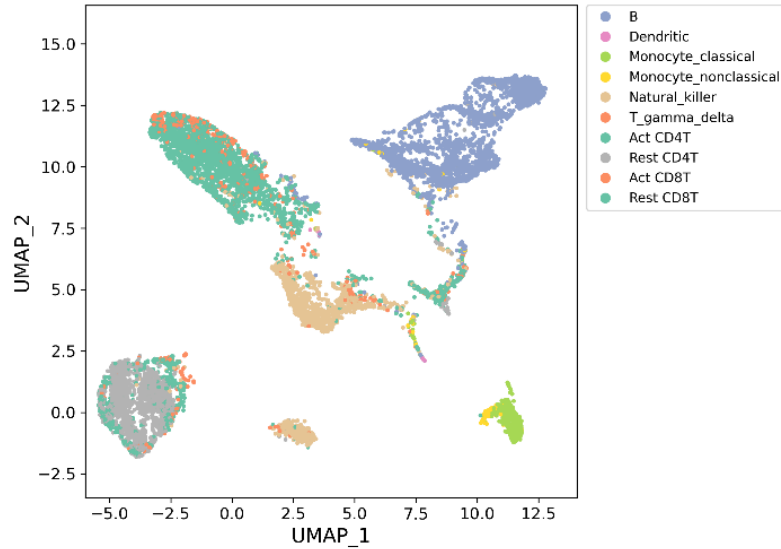


B.

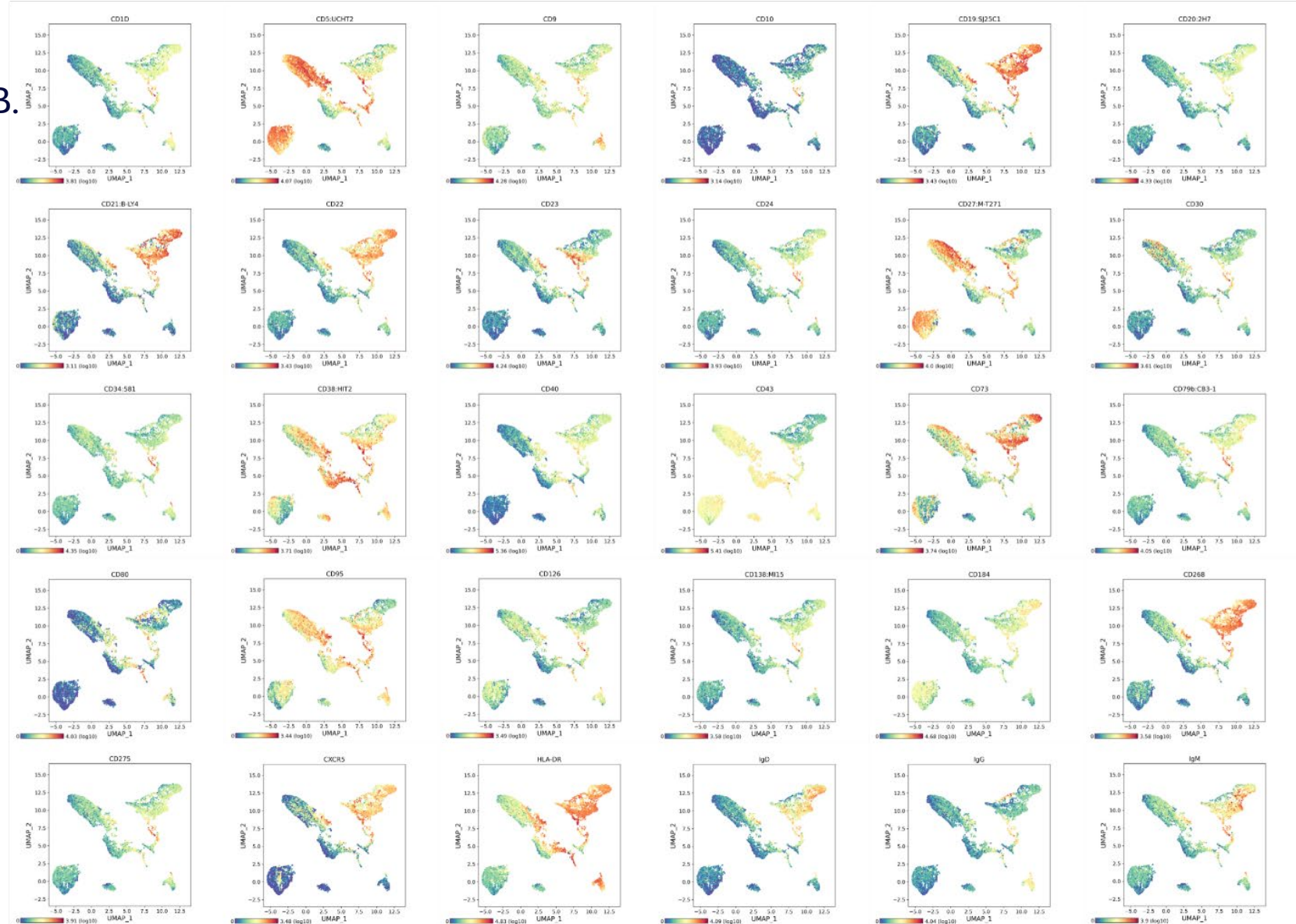


Detect 30 critical B-cell markers in your samples with confidence

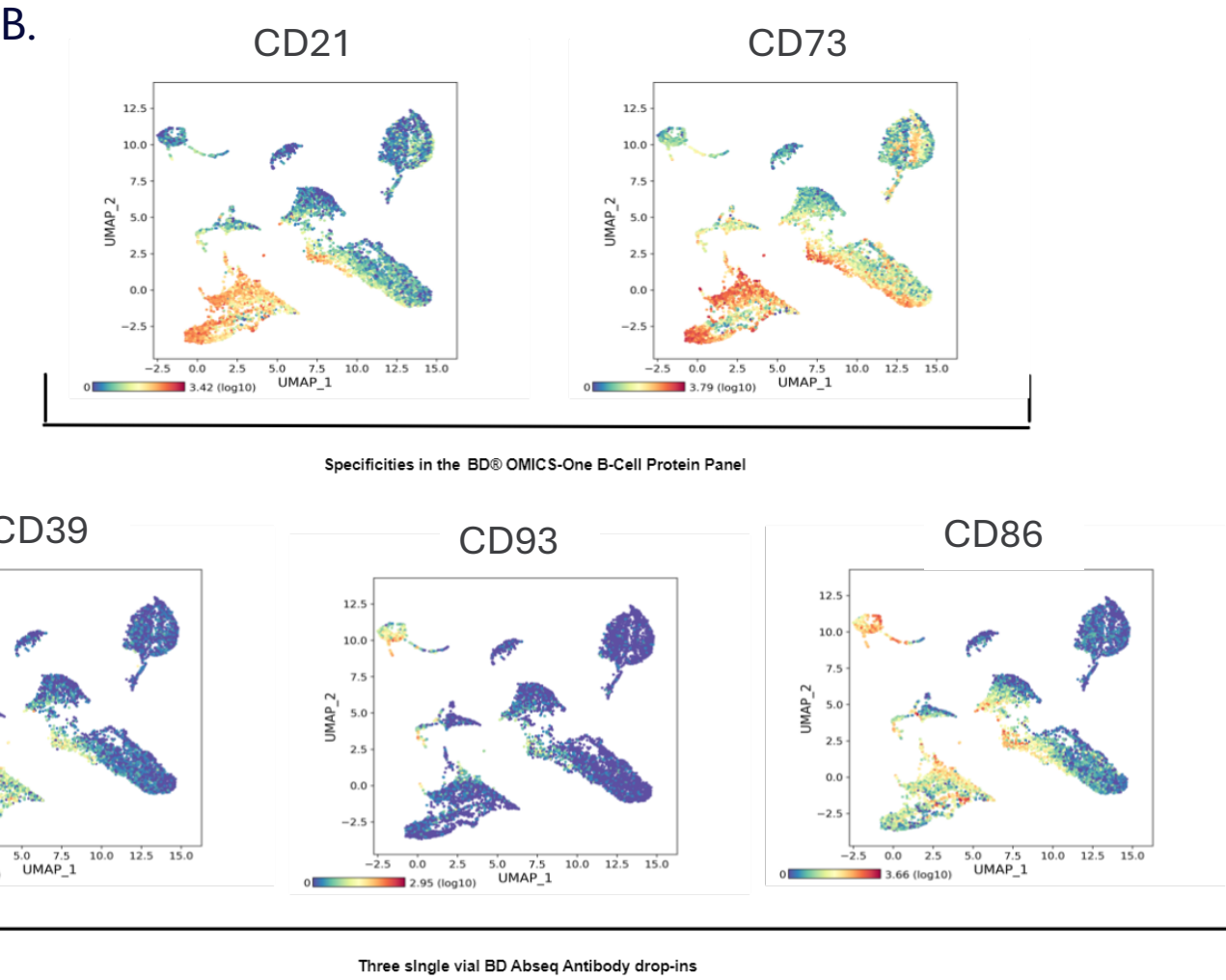
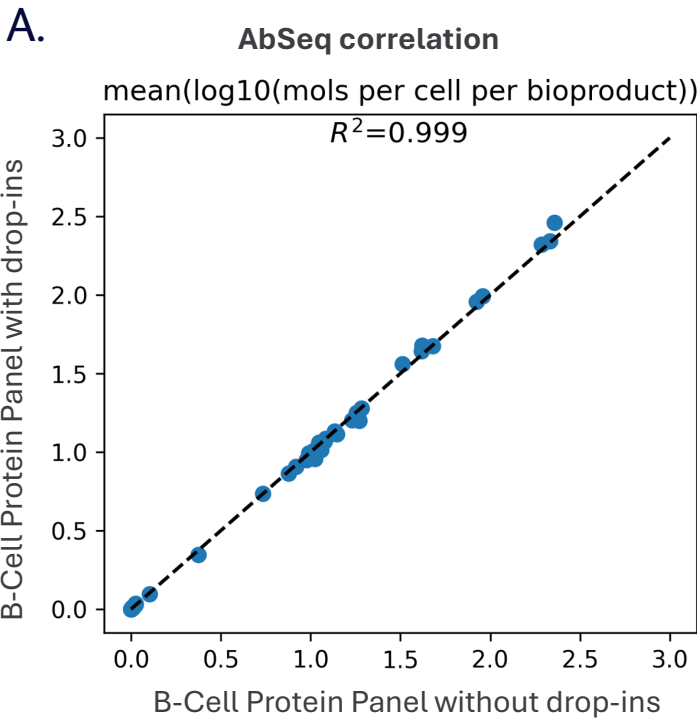
A.



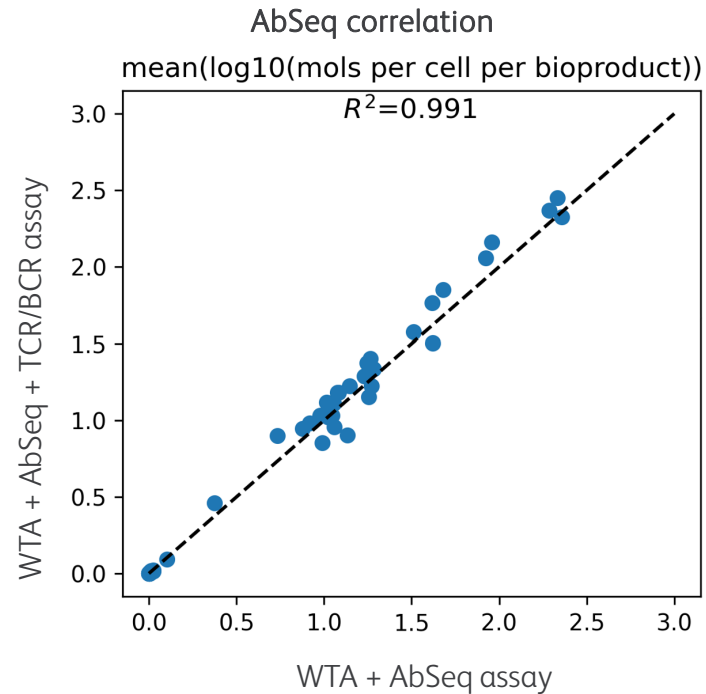
B.



Add additional specificities of interest to the BD® OMICS-One B-Cell Protein Panel without compromising performance

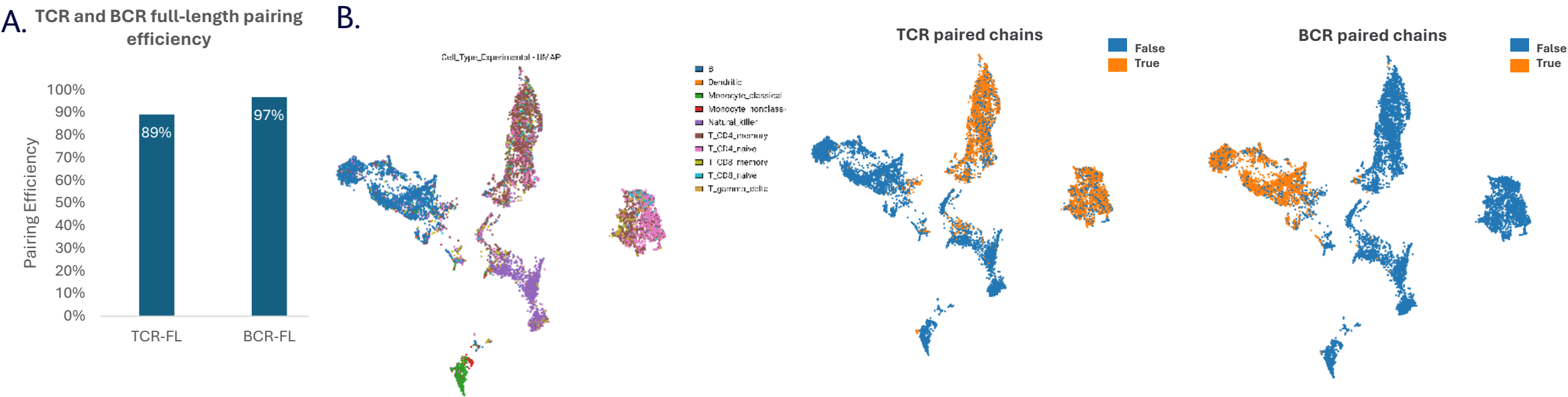


Multiomics enabled: BD[®] OMICS-One B-Cell Protein Panel is designed to work with WTA and TCR/BCR assays



The addition of BD Rhapsody™ TCR/BCR Next Assay does not impact BD[®] OMICS-One B-Cell Protein Panel performance

Multiomics enabled: BD[®] OMICS-One B-Cell Protein Panel is designed to work with WTA and TCR/BCR assays



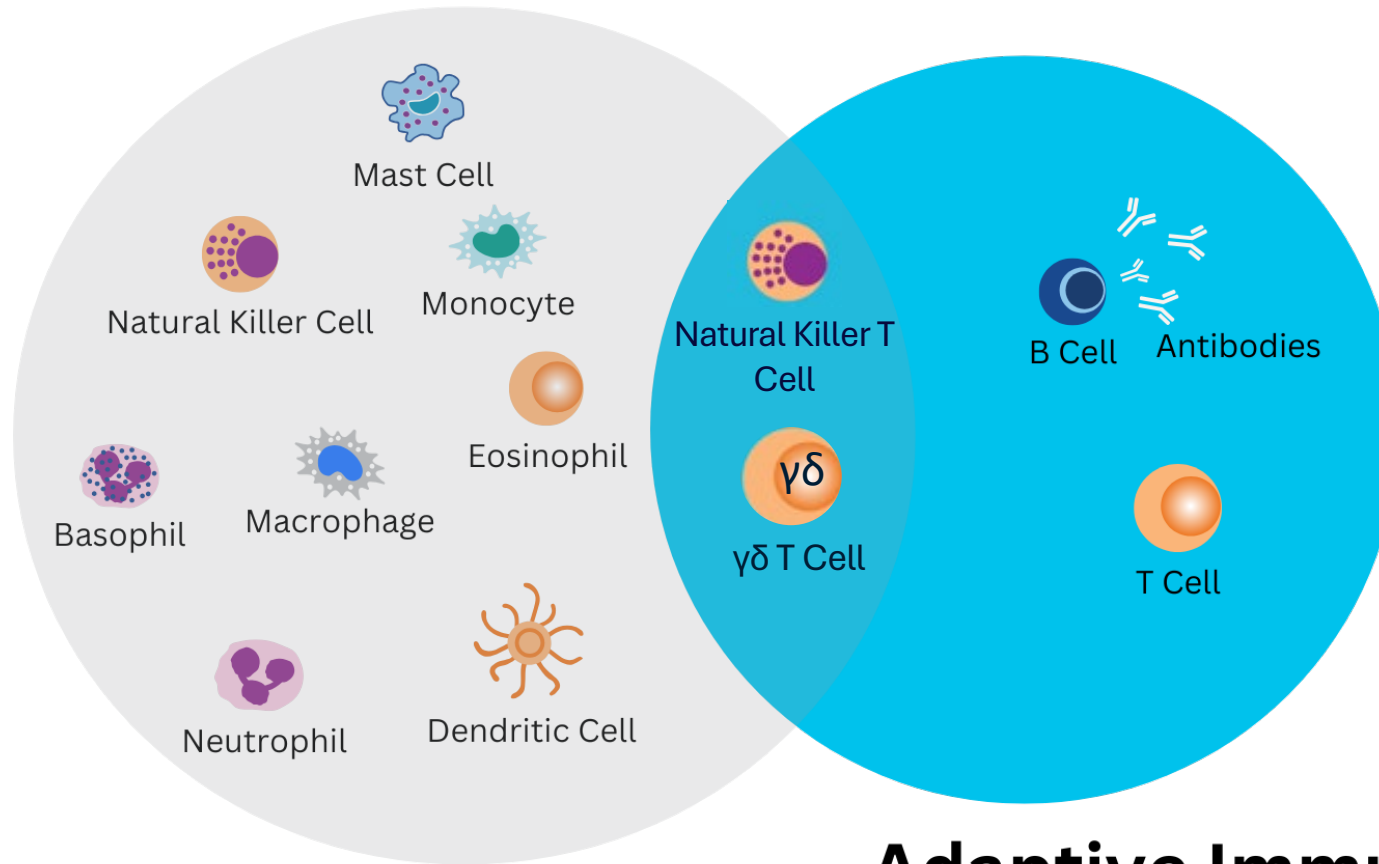
C. ● FWR1 ● CDR1 ● FWR2 ● CDR2 ● FWR3 ● CDR3 ● FWR4

Cell Index	BCR Chain	VDJ Translation Trimmed
1685	IG_Heavy	EVQLVESGGGLVQPGKSLRLSCTAS FTFGDY AMSWVRQAPGKLEWVG FIRSKAYG TT EYAASVKGRFTISRDDSKSIAYLQMNSLKTEDAVYYCTSSYYDFW SGYYPFDY WGQGLTV VSS
1685	IG_Kappa	DIQMTQSPSAMSASVGDRVTITCRAS QGISNYLAWFQ KPGKVPKRLIY AASLQSGVPSRFS SGSGSGTEFTLTIS SLQPEDFATYYCLQHNSYPWT FGQGTKVEIK
34654	IG_Heavy	EVQLVESGGGLVQPGKSLRLSCTAS FTFGDY AMSWVRQAPGKLEWV SAISGSGSTYYADSVKGRFTISRDN SKNTLYLQMNSLRAEDTAVYY CAKDQYSSGWTFDY WGQGLTVVSS
34654	IG_Lambda	SSELTQDPAVSVALGQTVRITCQGD SLRSYYASWYQ KPGQAPVLVIY GKNRPSGIPDRFSGSSSGNTASLTITGAQAEDEADYYCNSRDSSGNPVV FGGGTKLTVL
73399	IG_Heavy	QVQLVQSGAEVKKPGASVKVCKAS GYTFTSYDINWVRQATGQGLEWMGMNP NSGNTGYAQKFQGRVTMT RTSISTAYMELSSLRSEDTAVYYCARGGTWVG VVNPGGYYWGQGLTVVSS
73399	IG_Kappa	DIVMTQSPDSLAVSLGERATINCKSS QSVLYSSNNKNYLAWYQ KPGQPPKLLIYW ASTRESGVPDRFSGSGSGTDFTLTIS SLQAEDVAVYYC QQYYSTPYTFGQGTKLEIK
210497	IG_Heavy	EVQLVESGGGLVQPGKSLRLSCTAS FTFGDY AMSWVRQAPGKLEWV SAISGSGSTYYADSVKGRFTISRDN SKNTLYLQMNSLRAEDTAVYY CAKDYYDSSGYSKTHYYYYYGM DVWGQGT TVTVSS
210497	IG_Kappa	DIVMTQSPDSLAVSLGERATINCKSS QSVLYSSNNKNYLAWYQ KPGQPPKLLIYW ASTRESGVPDRFSGSGSGTDFTLTIS SLQAEDVAVYYC QQYYSTPRTFGQGTKVEIK
210530	IG_Heavy	QVQLQESGPGLVKPSQTLSTCTV SGSSISGSDYYW SWIRQPPGKLEWIG IYYSGSTYYNP SLKSRVTISVDTSKNQFSLKLSVTAADTAVYYC CAREAAVTYYGLDPPRWFD PWGQGLTVVSS
210530	IG_Kappa	DIQMTQSPSSLSASVGDRVTITCRAS QSISSYLNWYQ KPGKAPKLLIY AASLQSGVPSRFS SGSGSGTDFTLTIS SLQPEDFATYYCQSYSTPYTFGQGTKLEIK

BD[®] OMICS-One Adaptive Protein Panel

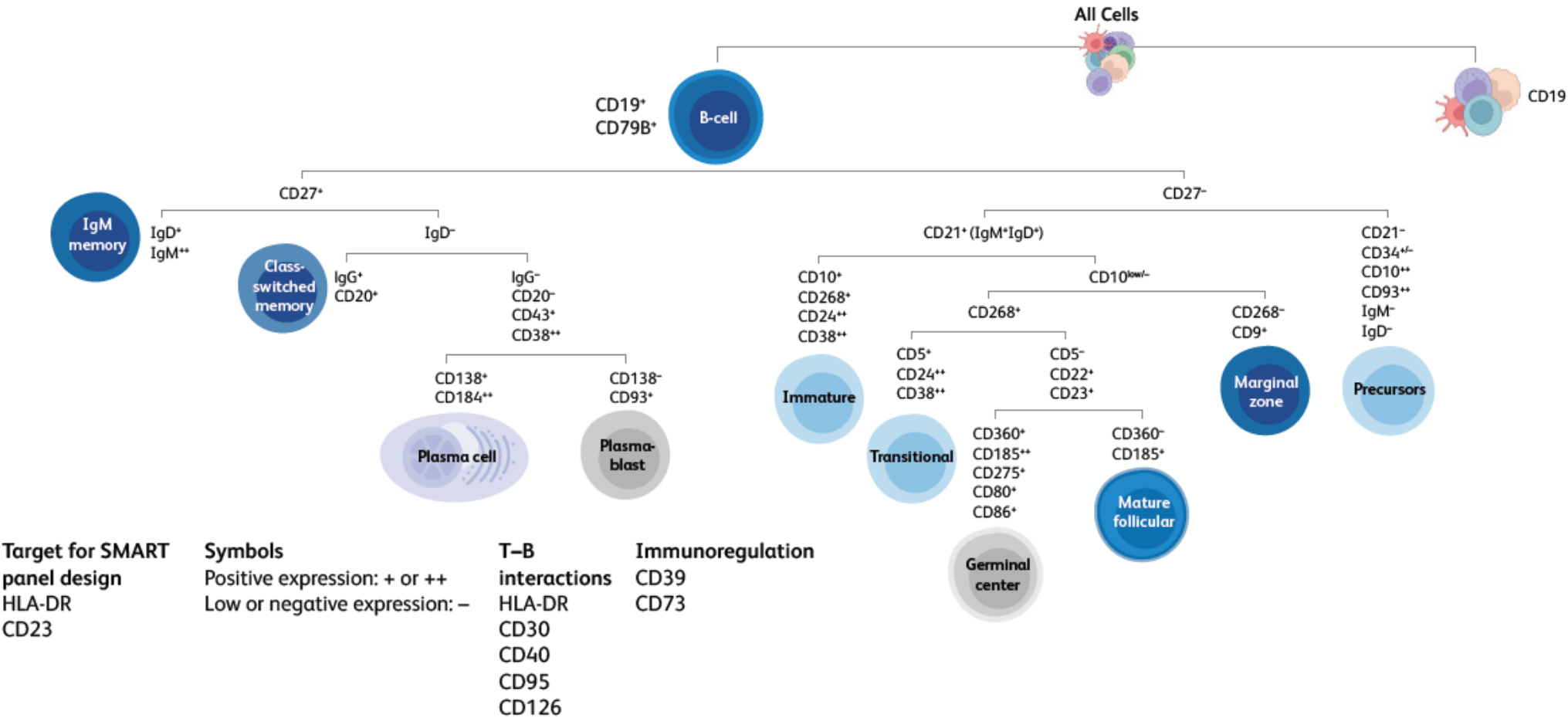
T-cells and B-cells are the two major players in adaptive immune system

Innate Immunity



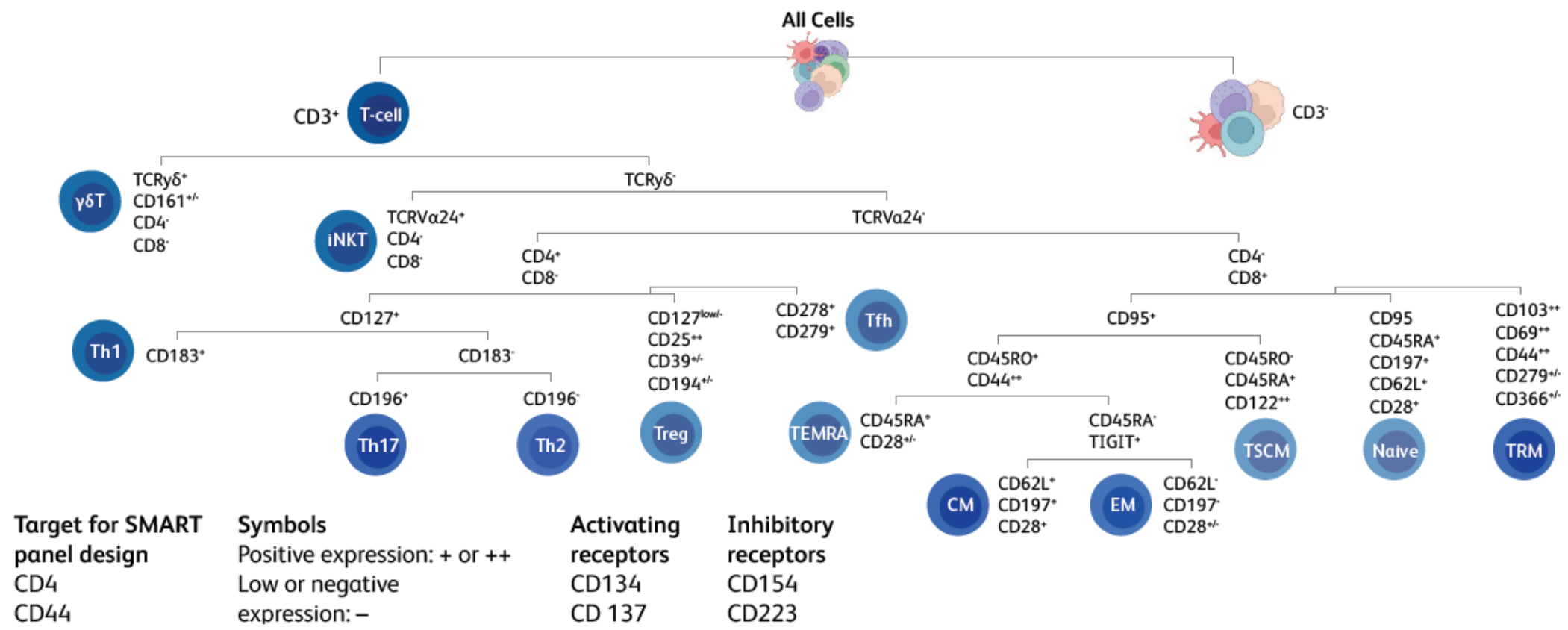
Adaptive Immunity

Explore B-cell populations and their activation and suppressed states



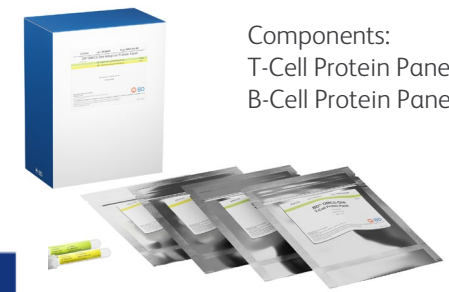
B-cell populations and activation states monitored by this panel.

Explore T-cell populations and their activation and suppressed states



T-cell populations and activation states monitored by this panel.

BD® OMICS-One Adaptive Protein Panel specificities



Components:
T-Cell Protein Panel (1 test/pouch, 2 pouches)
B-Cell Protein Panel (1 test/pouch, 2 pouches)

Specificity	Clone
CD1d	CD1d42
CD5	UCHT2
CD9	M-L13
CD10	HI10A
CD19	SJ25C1
CD20	2H7
CD21	B-LY4
CD22	HIB22
CD23	EBVCS-5
CD24	ML5
CD3	UCHT1
CD4*	SK3
CD8	SK1
CD25	2A3
CD28	L293
CD44*	L178
CD45RO	UCHL1
CD45RA	HI100
CD69	FN50
CD62L	DREG-56

Specificity	Clone
CD27	M-T271
CD30	BERH8
CD34	581
CD38	HB7
CD40	5C3
CD43*	1G10
CD73	AD2
CD79b	CB3-1
CD80	L307.4
CD95**	DX2
CD103	BER-ACT8
CD127	HIL-7R-M21
CD134	ACT35
CD137	4B4-1
CD154	TRAP1
CD161	HP-3G10
CD183	1C6/CXCR3
CD194	1G1
CD196	11A9
CD197	2-L1-A

Specificity	Clone
CD126	M5
CD138	MI15
CD184	12G5
CD185	RF8B2
CD268	11C1
CD275	2D3/B7-H2
HLA-DR*	G46-6
IgD	IA6-2
IgG	G18-145
IgM	G20-127
CD223	T47-530
CD272	J168-540
CD278	DX29
CD279	EH12.1
CD357	V27-580
CD366	7D3
TCRγ/δ	11F2
TCR Vα24-Jα18	6B11
TIGIT	tgMab-2

*Targets for SMART panel design

**The BD® OMICS-One Adaptive Protein Panel consists of two 30-plex protein panels: The BD® OMICS-One T-Cell Protein Panel and B-Cell Protein Panel. Both the T-cell and B-cell panel contain the same anti-CD95 antibody.

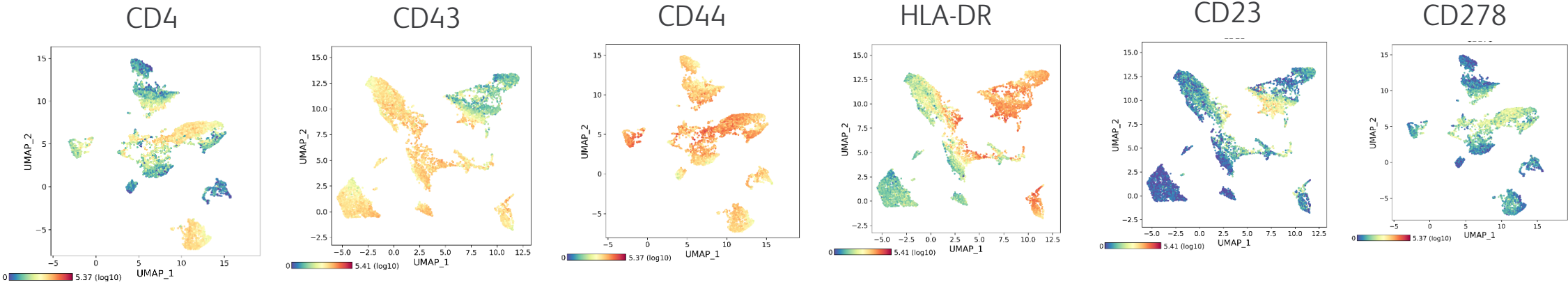
Fewer sequencing reads consumed by the high expressors and more reads allocated to lowly expressed markers

Percent of Total Sequencing Reads Consumed		
Markers	Without SMART panel design	With SMART panel design
Reduction of sequencing reads allocated to primary markers ▼		
CD44	23.20	5.34
HLA-DR	19.06	14.12
CD43	8.63	3.74
CD4	2.98	2.25
Read re-allocation to lowly expressed markers ▲		
CD3	1.16	2.58
CD62L	0.68	1.39
CD272	0.86	1.61
CD184	0.44	1.41
CD69	4.78	11.82
CD38	1.08	1.92
IgD	0.44	0.80
CD19	0.61	1.13
CD366	0.91	1.54
CD22	0.46	0.65

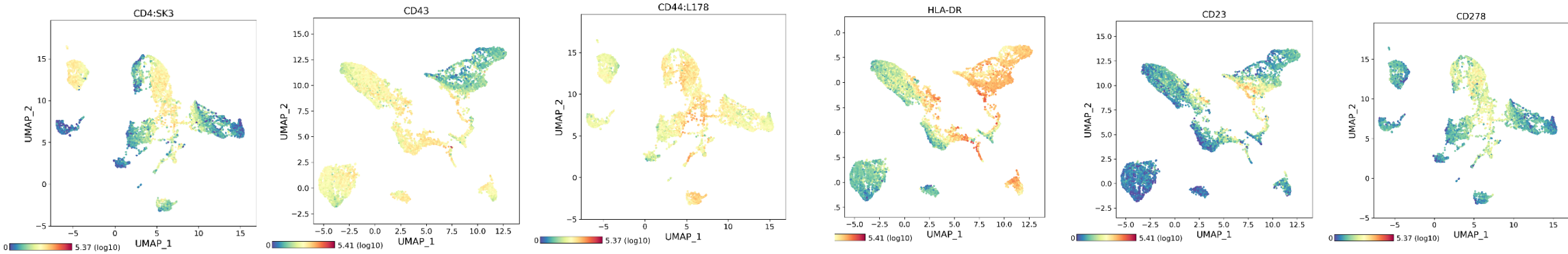
Percent of Total Sequencing Reads Consumed		
Markers	Without SMART panel design	With SMART panel design
CD197	0.46	1.38
IgG	0.33	0.44
CD40	0.47	0.91
CD185	0.19	0.48
CD25	1.87	4.34
CD1d	0.41	0.43
CD20	0.65	1.53
CD275	0.86	0.73
CD357	0.28	0.64
CD223	0.39	0.86
CD23	0.87	1.27
TIGIT	0.19	0.33
CD278	0.33	0.77
CD138	0.21	0.24
TCRγ/δ	0.17	0.33
CD154	0.07	0.14

CD4, CD43, CD44 and HLA-DR detection is not compromised, while better resolution of low expressors is found with SMART panel design

Without SMART panel design

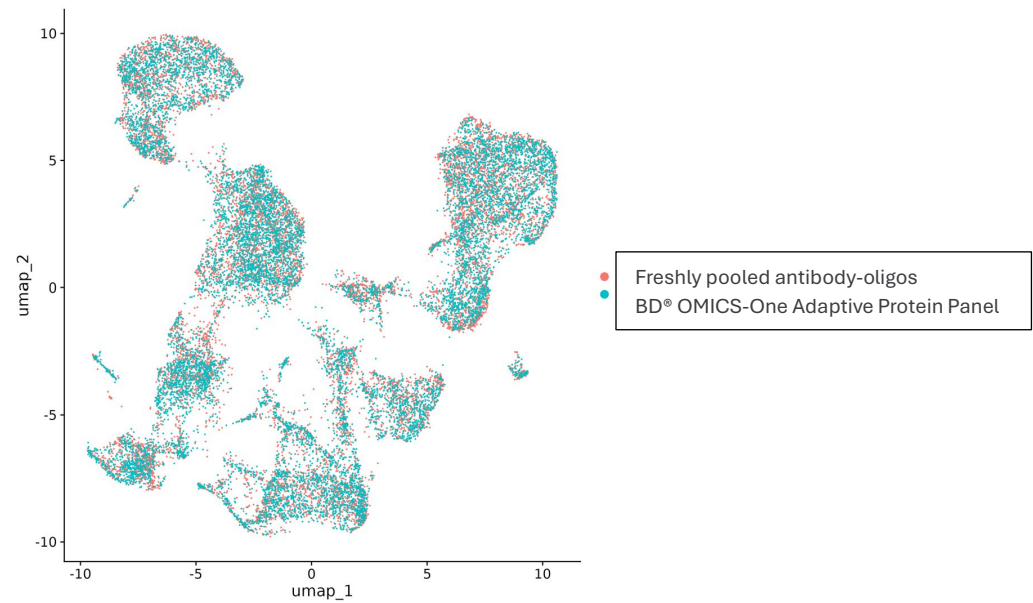


With SMART panel design

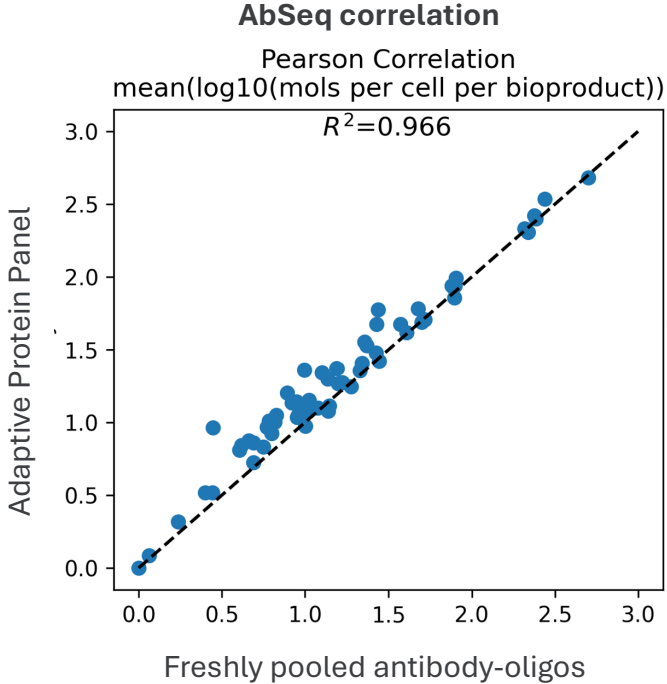


Comparable performance of the lyophilized panel to freshly pooled BD[®] AbSeq Antibody-Oligo Reagents

A.

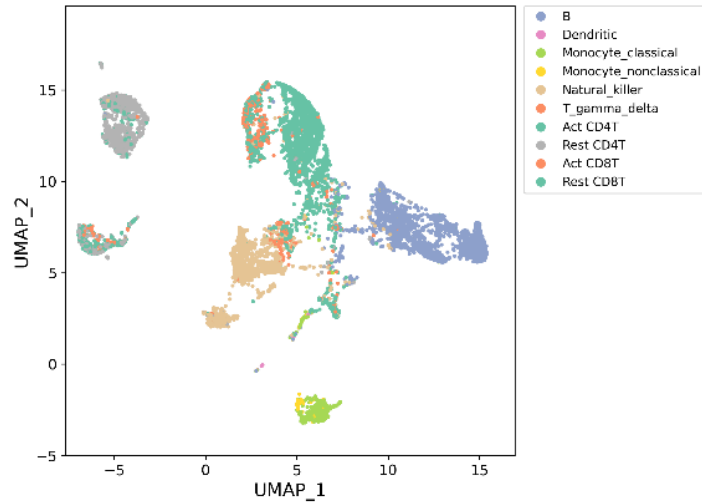


B.

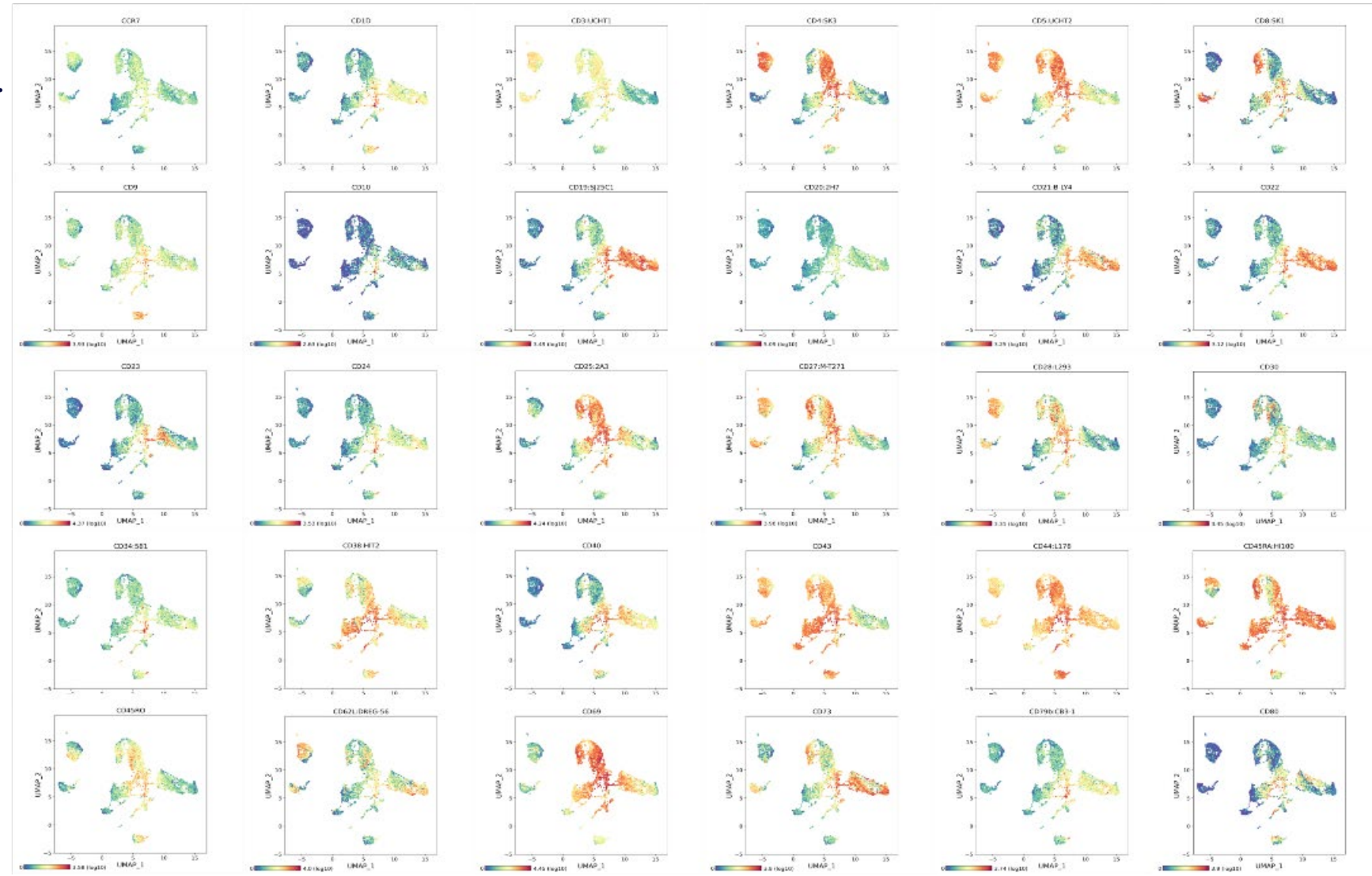


Detect 59 critical adaptive markers in your samples with confidence

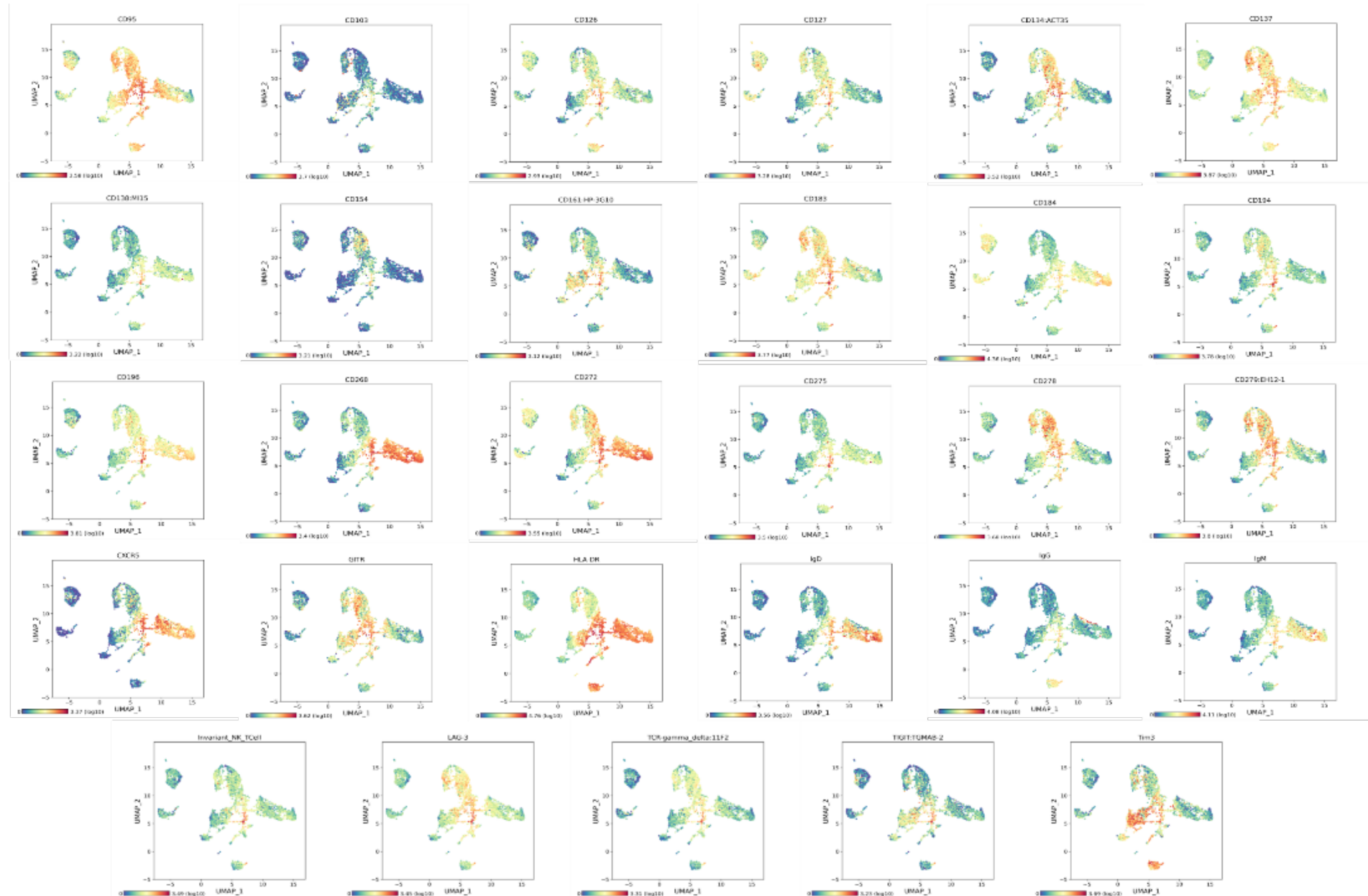
A.



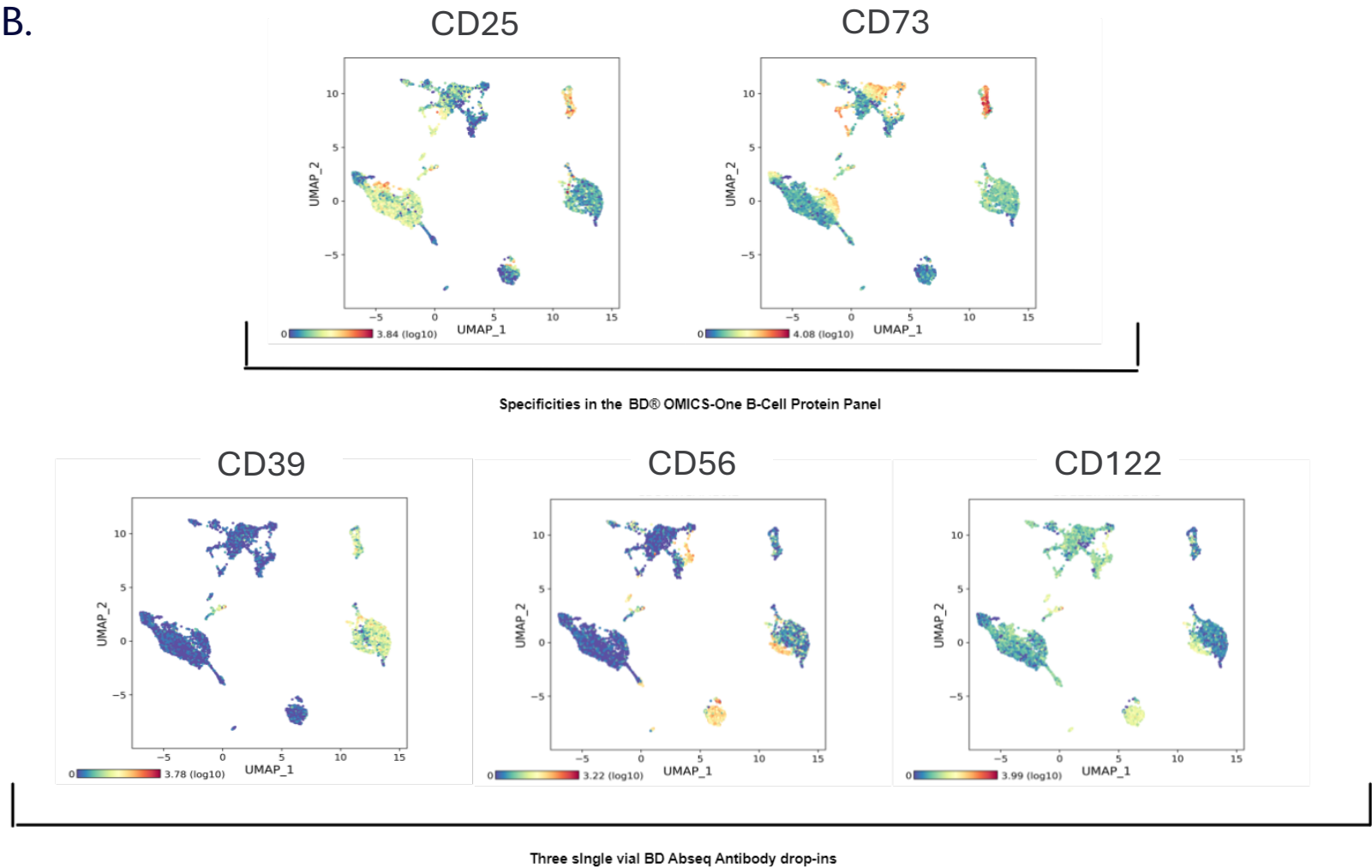
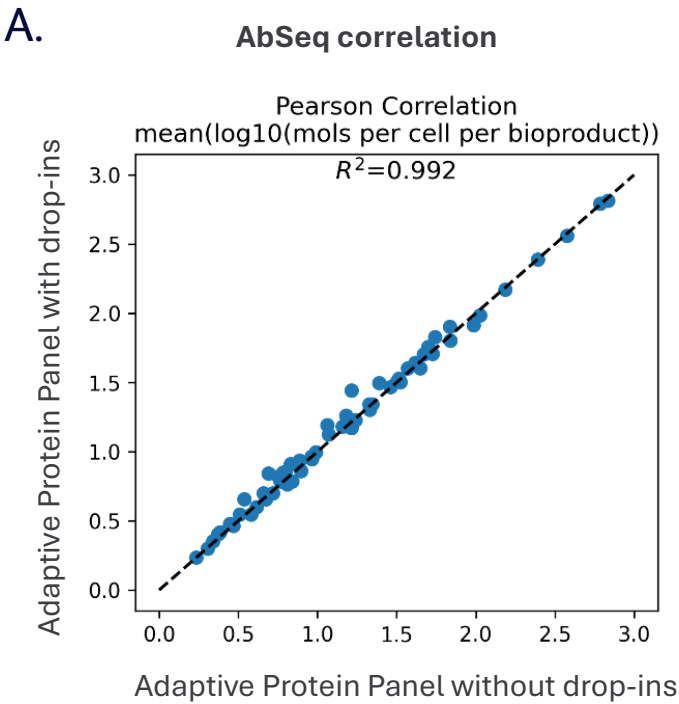
B.



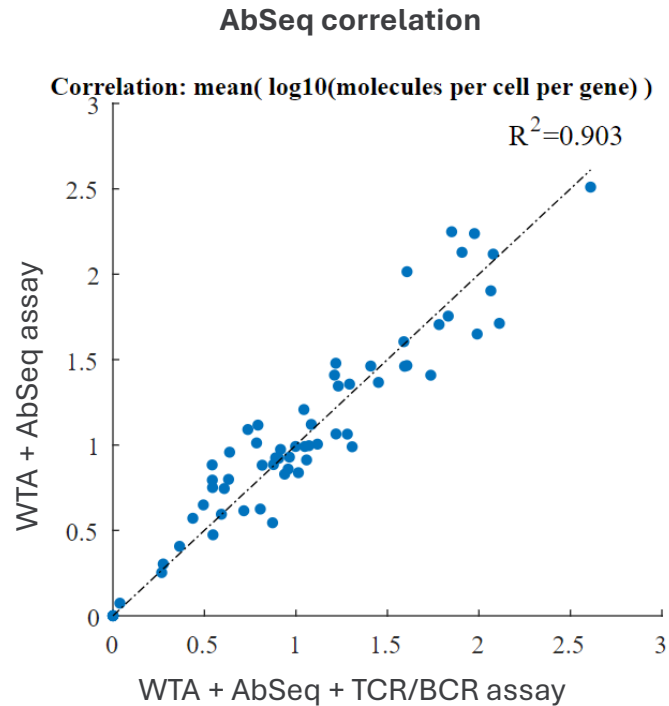
B. (cont.)



Add additional specificities of interest to the BD® OMICS-One Adaptive Protein Panel without compromising performance

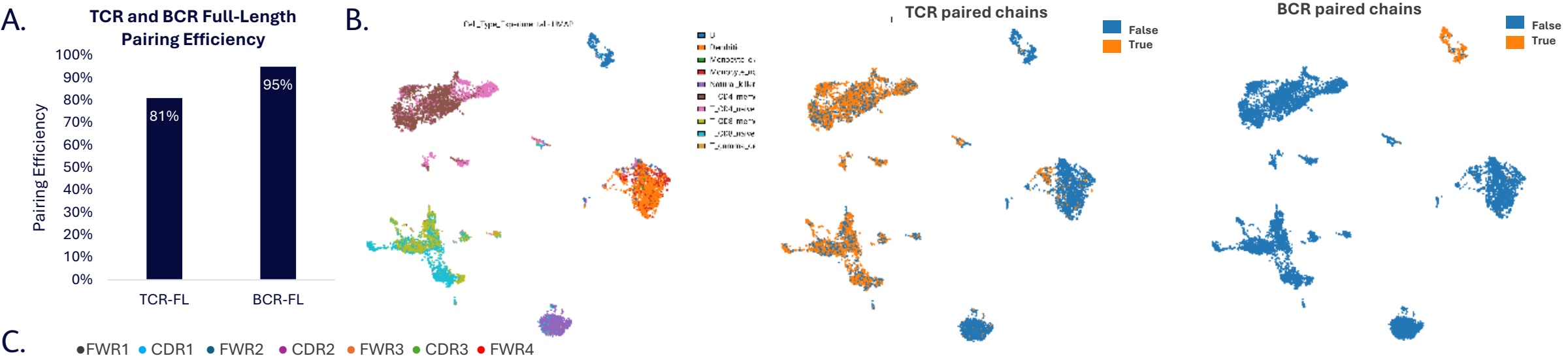


Multiomics enabled: The BD[®] OMICS-One Adaptive Protein Panel is designed to work with whole transcriptome and TCR/BCR assays



The addition of a TCR/BCR profiling assay does not impact BD[®] OMICS-One Adaptive Protein Panel performance

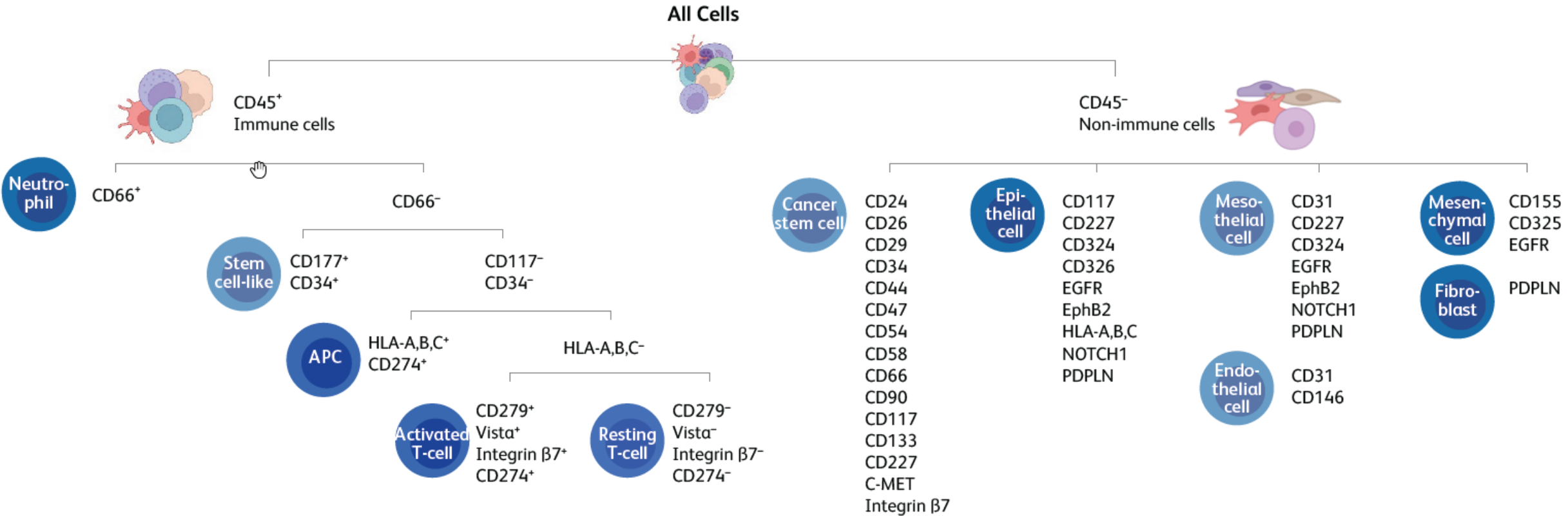
Multomics enabled: The BD[®] OMICS-One Adaptive Protein Panel is designed to work with whole transcriptome and TCR/BCR assays



Cell Index	TCR/BCR Chain	VDJ Translation Trimmed
23120	TCR_Alpha	AQSVSQHNHHVILSEAAASLELGCNYS GGTVN LFWVYQPGQHLQLLLKY FS G DPL V KG IKGF EA E FI KS FS FNLRKPSVQWSDTA EY FC AV KEGT NA G K ST FG DG TTL TVKP
23120	TCR_Beta	EAQVTQNPRYLITVTGKKLTVCQSN MN HEYMSWYRQDPGLGR QI Y YS MN VE VD KGDVPEGYKVS R KEKRN FL ILESPSPNQTS LY FCASW GL T GE LF FG EGS R LT VL
33061	TCR_Alpha	AQSVTQLDSHVSSEGTPVLLRCNYS SS Y SP SLFWYVQHPNKG LQ LL LY TS AA TLV KG ING FE A EF K S KT SF HLTKPSAHMSDAA EY FC V SPGYSS AS K II IF GS G TRLS IR P
33061	TCR_Beta	NAGVTQTPKFRVLKTGQSM TL CAQD MN HEYMYWYRQDPGMGLRL I HY SV GE GT TAKGEVPDGYNV S RLKKQ N FL LG ESAAPSQTS VY FCASSY SM DRGSD TQ Y FG PG TR LL TV L
39942	TCR_Alpha	QQQVKQSPQSLIVQKGGIS I NCAYE NT AFD Y FPWYQ Q FPKG P ALL IA IRPD VS E KE GRFT IS FNK SA Q FS LHIMDSQPGDSAT Y CA AS KGS NY QL I WGAG T KL IK P
39942	TCR_Beta	GAVVSQHPSWVICKSGTSVKIECRSL DF Q AT TMFWYRQ FP KQSLML MA TS NE G SK AT Y EQGVEKD K FLINHASLTLSTLTVTSAHPEDSS FY ICSA II PEET QY FG PG TR LL VL
59553	IG_Heavy	EVQLVESGGGLVQPGGSLKLS CA AS GF T ST YW M H W VRQSPGKGLEW V SR INT G ES HT NY AD IS YAD F VEGRFTTSRDNAKNTLHLQM HS LRAEDSGV Y YCARALE Y FYGLD W WG QT TV IV SS
59553	IG_Kappa	DIVMTQSPDSLAVSLGERATINCKSS Q SVLYSS K N KN YAWYQ Q KPGQPPN LI Y W AST RE SGVPDR FS GS G SGTN FT L IS RLQAEDVAV Y Y C Q Y Y S PP T FG GG T K VE IK
65072	IG_Heavy	EVQLVESGGGLVQPGRLSL S CTAS GF T FG DY A MSWVRQAPGKGLEW V GF IR SKAY G GT TE YAASV K GRFTISRDDSK SI AYLQMNSL K TEDTAV Y Y C TS SV SWWY T ALD AF DI W G Q GT M TV SS
65072	IG_Kappa	DIQMTQSPSSLSASVGD R VTITCRAS Q GIS NY LAWYQ Q KPGK V PK LI Y AA STL Q SGVP S RFSGSGSGTD FT L TI SS LQ PE D VAT Y Y C K Y NSAP IT FG Q G TR LE IK
266242	IG_Heavy	EVQLVESGGGLVQPGGSLRL S CAAS GF T SS Y EM NWVRQAPGKGLEW I SY IS T SG ST Y SADSVRGRFTISRDN AK NSQY L QMNSLRAEDTAV Y Y C ARRG IS T IG FG AF DI W GRGT M TV SS
266242	IG_Kappa	DIVLTQTPPSLSVTPGQPASISCKSS Q SL LH IDG K TYLYWY L QKPGQSP QL LI Y EV SN RFSGVPDR FS GS G SGTD FT L K ISRVEAEDVG V Y C L Q TI HL PAL S FG GG T K VE IK

BD[®] OMICS-One Tumor Protein Panel

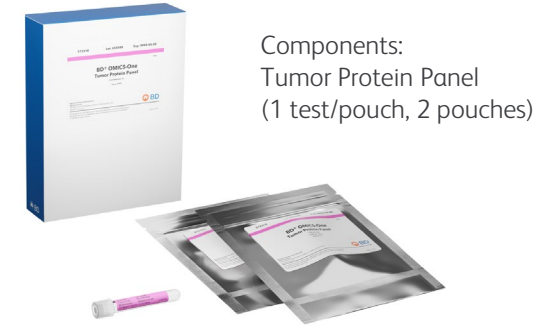
Explore tumor biology, immune evasion and stemness of your samples



Target for SMART panel design
CD44, CD45, HLA-A,B,C

Symbols
Positive expression: + or ++
Low or negative expression: -

BD® OMICS-One Tumor Protein Panel specificities



Specificity	Clone
CD24	ML5
CD26	M-A261
CD29	MAR4
CD31 (PECAM-1)	WM59
CD34	581
CD44*	L178
CD45*	HI30
CD47	B6H12
CD54	HA58
CD58	1C3

*SMART-titrated targets

Specificity	Clone
CD117	YB5.B8
CD133	W6B3C1
CD146	P1H12
CD155	TX24
CD227 (MUC1)	HMFG2
CD66	B1.1/CD66
CD90	5E10
CD274 (PD-L1)	MIH1
CD279 (PD-1)	EH12.1
CD324 (E-Cadherin)	67A4

Specificity	Clone
CD325 (N-Cadherin)	8C11
CD326 (EpCAM)	EBA-1
c-MET	3D6
EGFR	EGFR.1
EphB2	2H9
HLA-A,B,C*	G46-2.6
Integrin β 7	FIB504
Notch1	MHN1-519
Podoplanin	LpMab-17
Vista	MIH65.rMAb

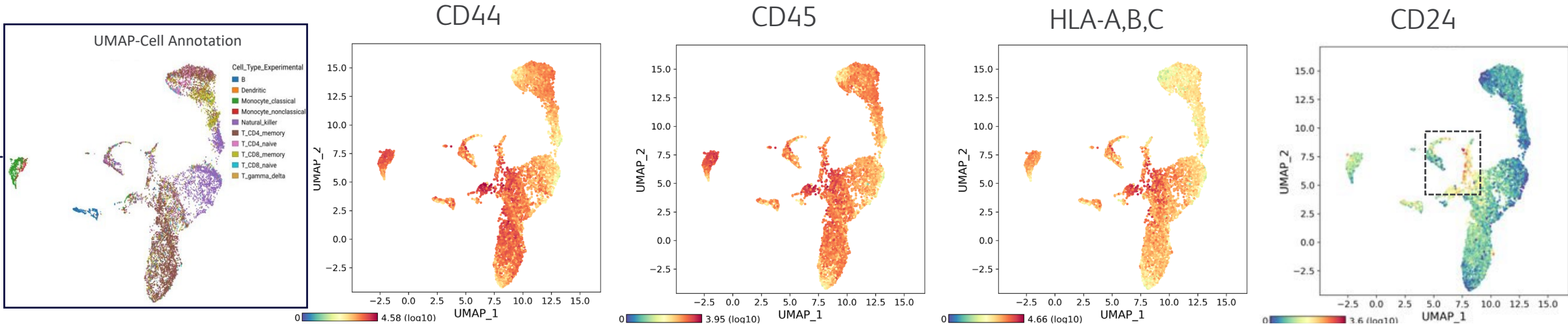
Fewer sequencing reads consumed by CD44, CD45 and HLA-A,B,C and more reads allocated to lowly expressed markers

Percent of Total Sequencing Reads Consumed		
Markers	Without SMART panel design	With SMART panel design
Reduction of sequencing reads allocated to primary markers ▼		
CD44	35.47	19.23
HLA-A,B,C	18.41	15.14
CD45	10.48	5.03
Read re-allocation to lowly expressed markers ▲		
CD54	7.83	14.72
CD47	4.58	4.68
CD34	3.46	4.13
EGFR	2.04	3.64
CD58	2.02	3.00
CD66	1.75	2.70
CD279	1.65	2.69
Podoplanin	1.24	2.54
CD155	1.13	2.04
CD24	1.00	2.04
CD29	0.84	1.78

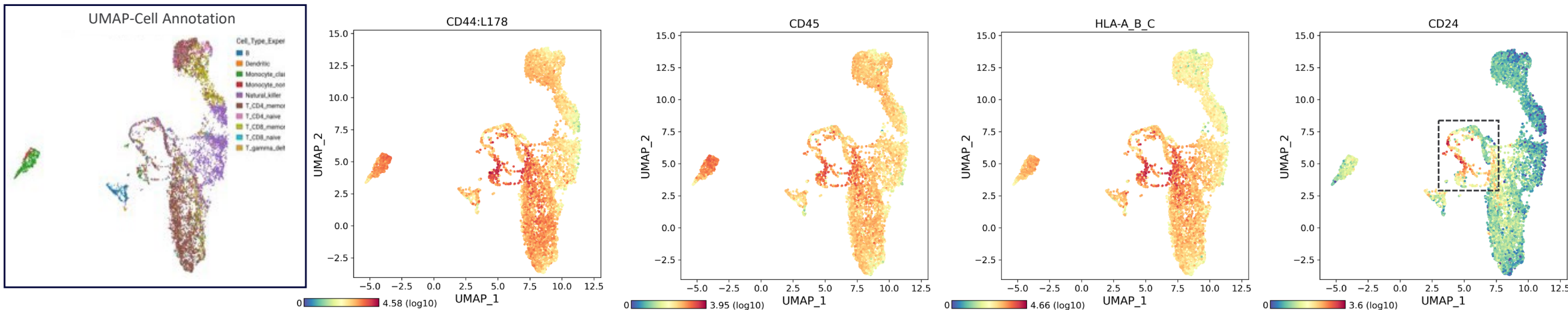
Percent of Total Sequencing Reads Consumed		
Markers	Without SMART panel design	With SMART panel design
CD31	0.84	1.70
CD274	0.79	1.67
CD117	0.78	1.62
CD26	0.75	1.58
CD324	0.73	1.54
Notch1	0.73	1.46
CD227	0.66	1.24
EphB2	0.52	1.15
CD90	0.47	0.96
Integrin β 7	0.43	0.90
c-MET	0.37	0.70
CD326	0.32	0.67
Vista	0.30	0.60
CD133	0.20	0.38
CD325	0.19	0.37
CD146	0.04	0.13

CD44, CD45 and HLA-A,B,C detection is not compromised, while better resolution of low expressors is found with SMART panel design

Without SMART panel design

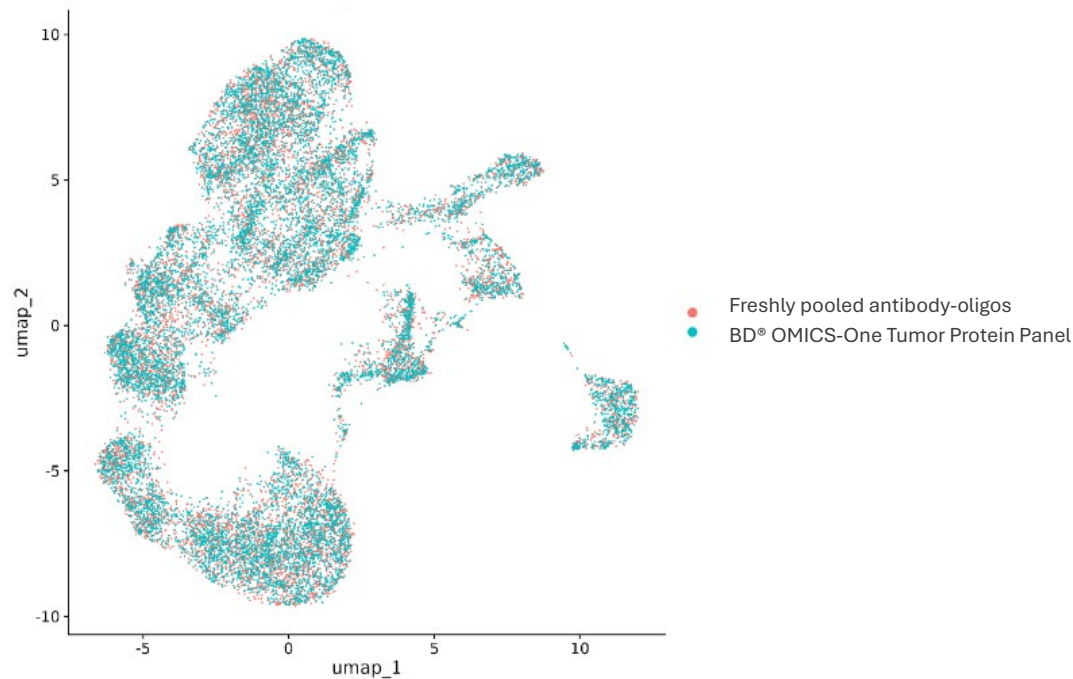


With SMART panel design

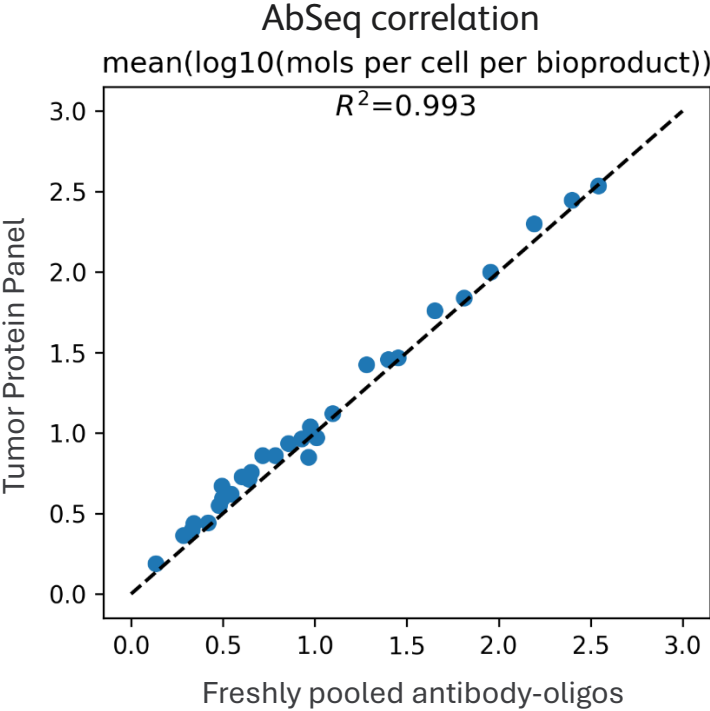


Comparable performance of the lyophilized panel to freshly pooled BD[®] AbSeq Antibody-Oligo Reagents

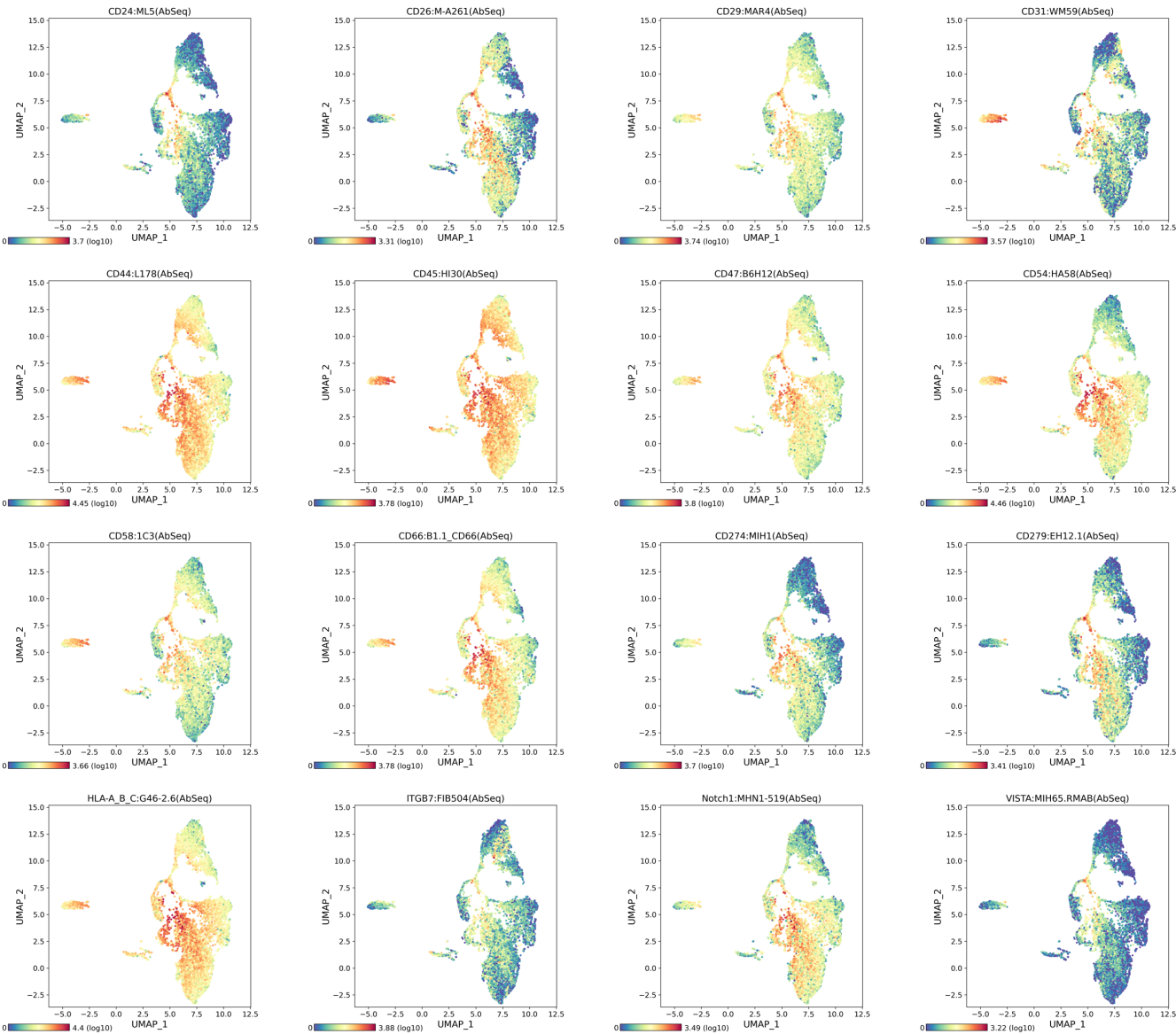
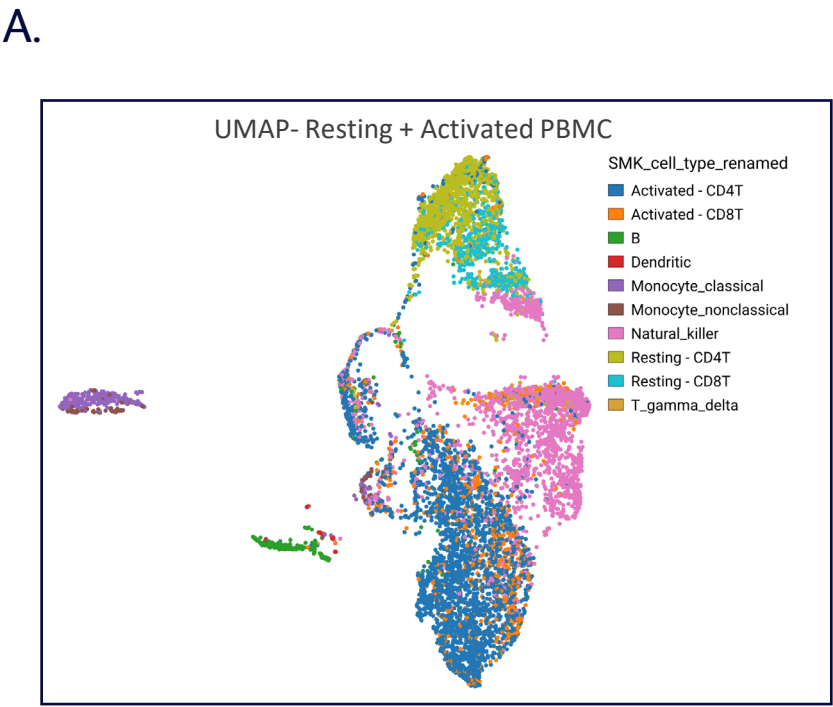
A.



B.

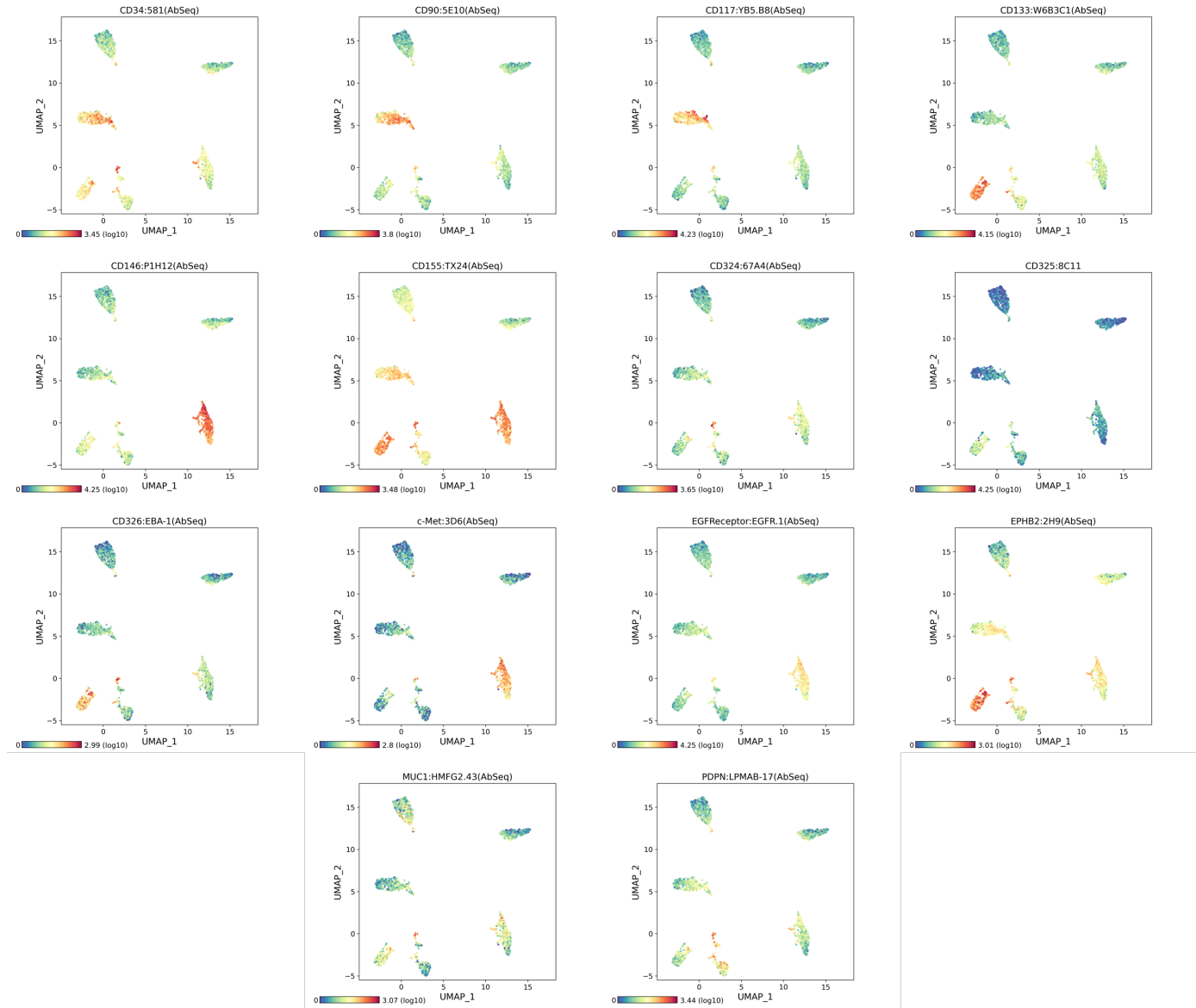
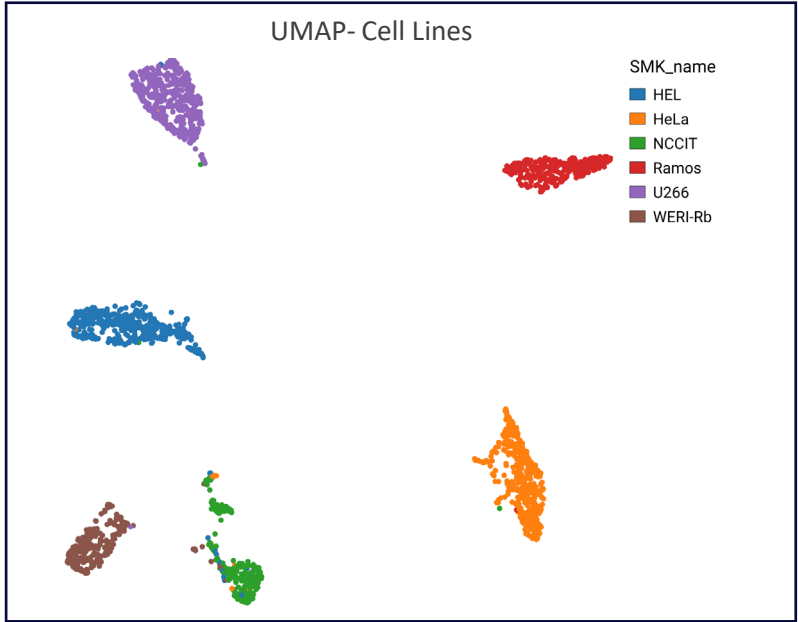


Detect 30 critical tumor markers in your samples with confidence

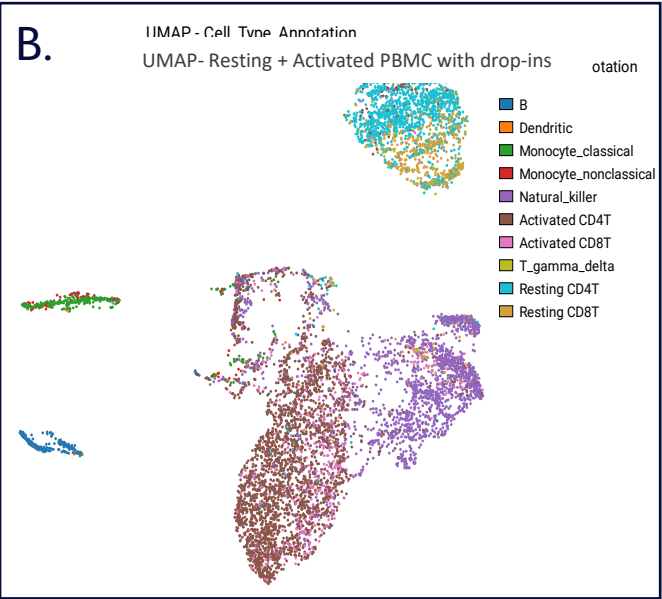
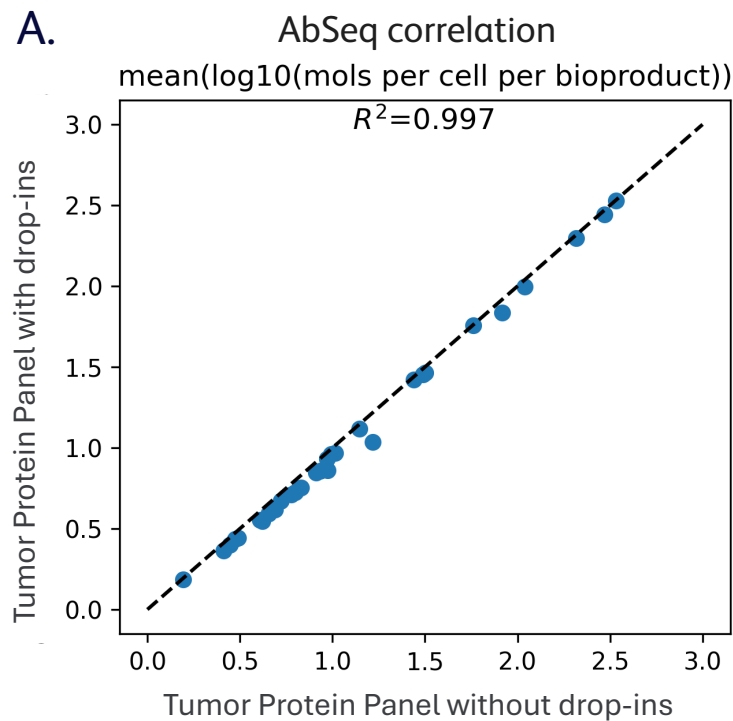


Detect 30 critical tumor markers in your samples with confidence

B.

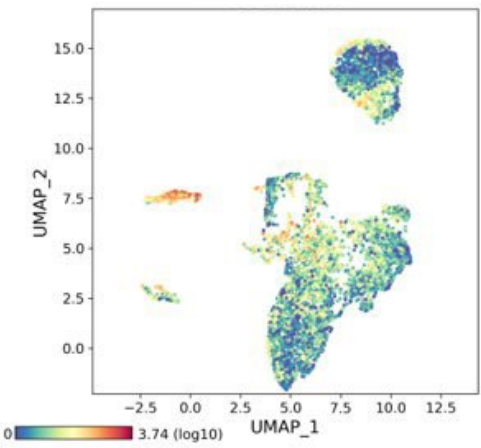


Add additional specificities of interest to the BD® OMICS-One Tumor Protein Panel without compromising performance

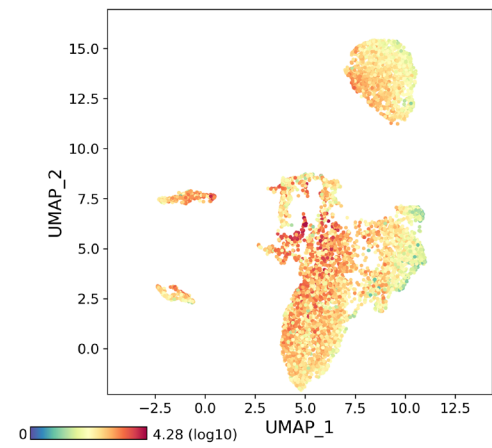


Specificities in the BD® OMICS-One Tumor Protein Panel

CD31

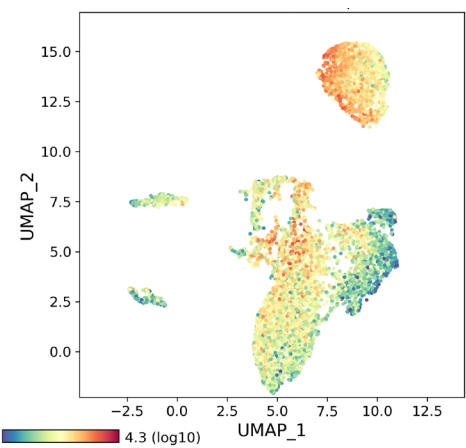


CD44

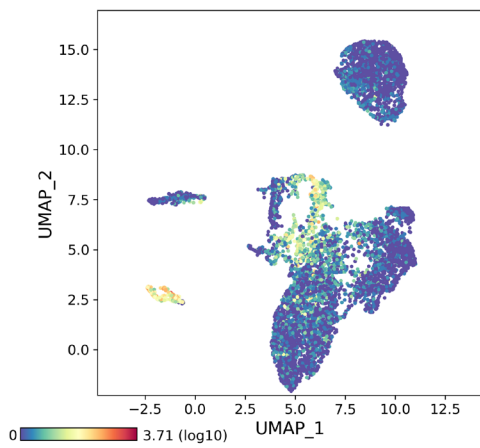


Three single-vial BD® AbSeq Antibody-Oligo drop-ins

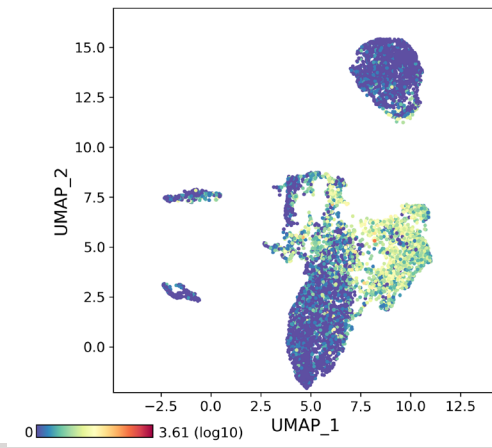
CD3



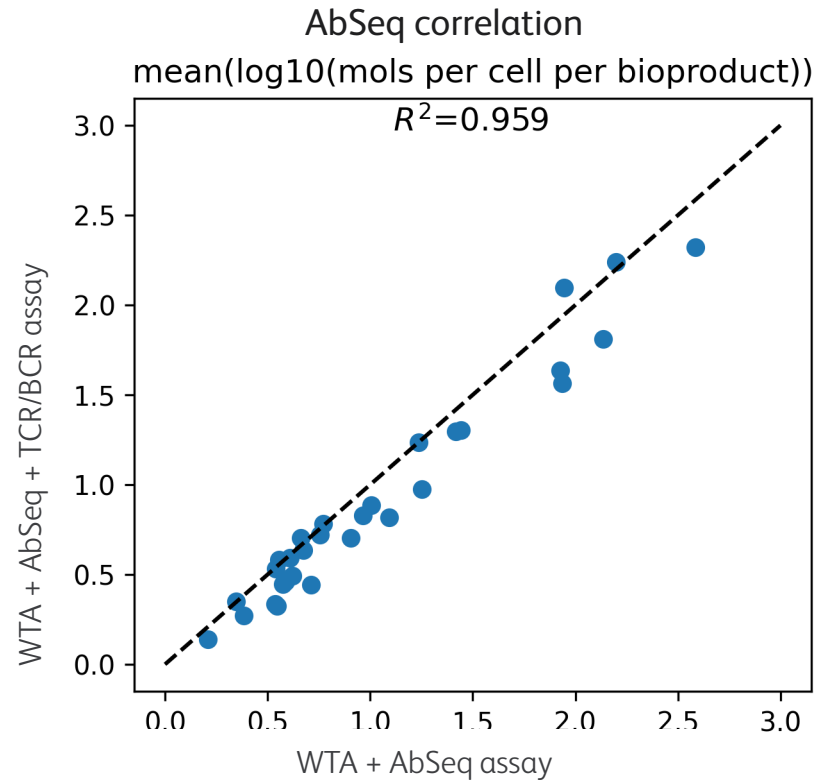
CD19



CD56

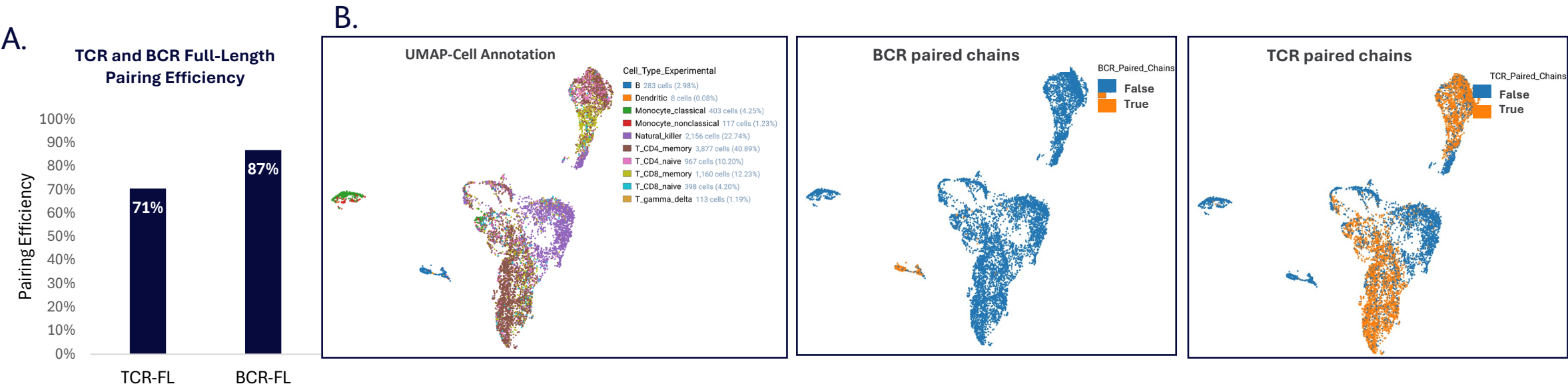


Multiomics enabled: BD[®] OMICS-One Tumor Protein Panel is designed to work with WTA and TCR/BCR assays



The addition of BD Rhapsody[™] TCR/BCR Next Assay does not impact BD[®] OMICS-One Tumor Protein Panel performance

Multiomics enabled: BD® OMICS-One Tumor Protein Panel is designed to work with WTA and TCR/BCR assays

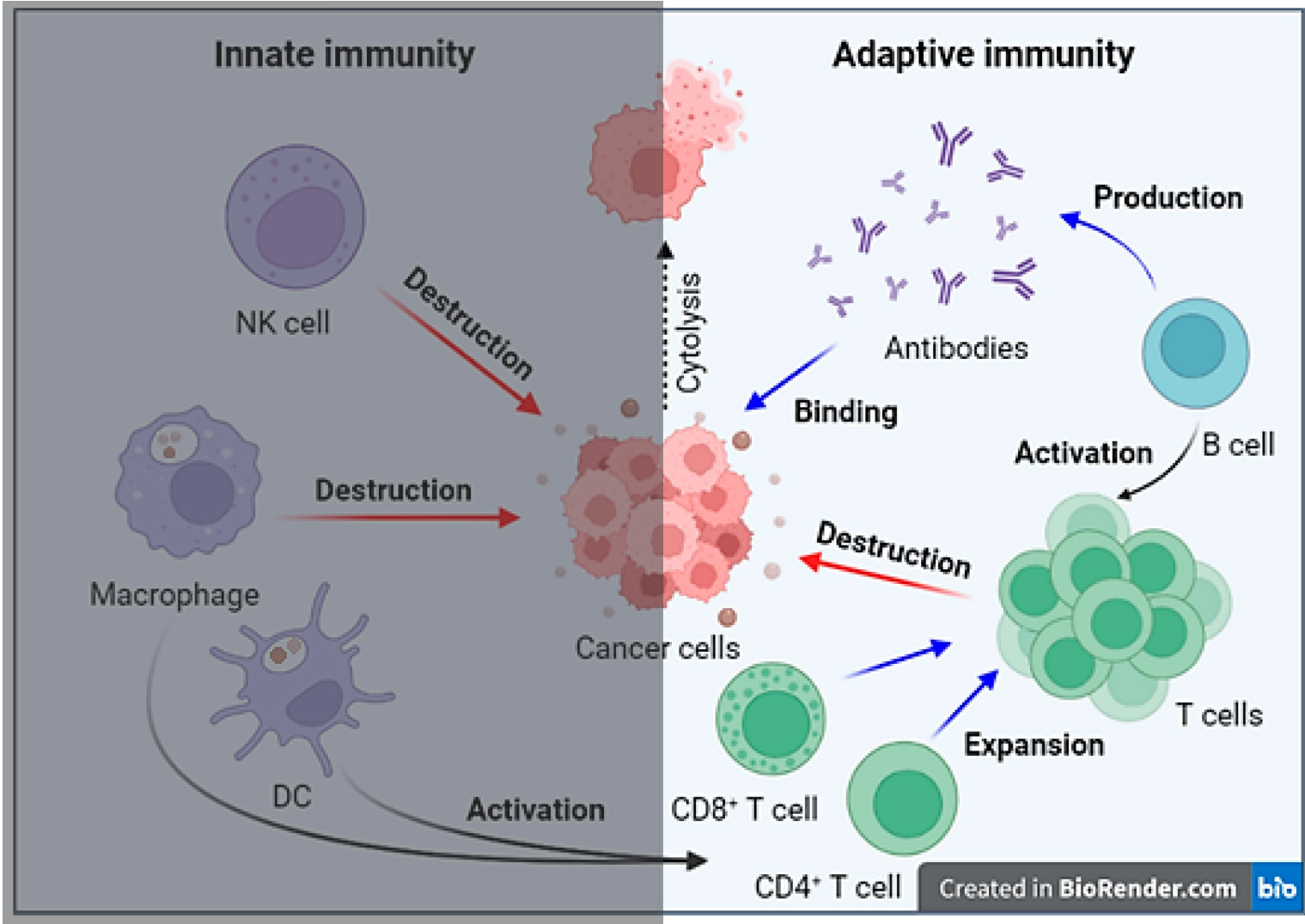


C. ● FWR1 ● CDR1 ● FWR2 ● CDR2 ● FWR3 ● CDR3 ● FWR4

Cell Index	TCR/BCR Chain	VDJ Translation Trimmed
15598	TCR_Alpha	VTSLSAYSPILIVQKGGIISIINCAYE NTAFDY FPWYQQFPGKGPALLIA IRPDVSE KKEGRFTISFNKSAKQFSLHIMDSQPGDSATYFC AARNTGGFKTI FGAGTRLFVKA
15598	TCR_Beta	AAGVIQSPRHLIKEKRETATLKCYPI PRHDT VYWYQQGPGQDPQFLIS FYEKMQ SDKGSIPDRFSAQQFSDYHSELNMSSLELGDSALYFC ASSQTGHGDTDTQY FGPGTRRLTVL
17194	TCR_Alpha	DQQVKQNSPQSVQEGRISILNCDYT NSMFDY FLWYKKYPAEGPTFLIS ISSIKDK NEDGRFTVFLNLSAKHLSLHIVPSQPGDSAVYFC AAHNAGNMLT FGGGTRLMVKP
17194	TCR_Beta	KAGVTQTPRYLIKTRGQVTLSCSPI SGHRS VSWYQQTPGQGLQFLFE YFSETQ RNKGNGPGRFSGRQFSNSRSEMNSTLELGDSALYLC ASSLRTGLRGGTDQY FGPGTRRLTVL
22191	TCR_Alpha	DQQVKQNSPQSVQEGRISILNCDYT NSMFDY FLWYKKYPAEGPTFLIS ISSIKDK NEDGRFTVFLNLSAKHLSLHIVPSQPGDSAVYFC AATFYGGATNKLI FGTGTLAVQP
22191	TCR_Beta	NAGVTQTPKFRVLKTGQSMTLCAQD MNHEY MYWYRQDPGMGLRLIHY SVGEGT TAKGEVPDGYNVSRLLKQNFLLGLESAAQTSVYFC ASSYSRQTQY FGPGTRRLTVL
23264	IG_Heavy	EVQLVESGGGLVKPGGSLRLSAAAS GFTFSYS MNWVRQAPGKGLEWVSS ISSSSSYI YYADSVKGRFTISRDNAKNSLYLQMNSLRAEDTAVYIC AREKVYCSGGSCYSLGAFDI WGQGTMTVTVSS
23264	IG_Kappa	AIRMTQSPSSFASSTGDRVTITCRAS QGISSY LAWYQKPGKAPKLLIY AAS TLQSGVPSRFSGSGSGTEFTLTISLQPDGFATYYC QQYNLSLR SAKGPRWKS
31565	TCR_Alpha	KQEVTTQIPAALSVPEGENLVNCSFT DSAIYN LQWFRQDPGKGLTSLLL IQSSQRE QTSGRNLNASLKDSSGRSTLYIAASQPGDSATYLC AVSKVAGNQFY FGTGTSLTVIP
31565	TCR_Beta	NAGVTQTPKFRVLKTGQSMTLCAQD MNHEY MYWYRQDPGMGLRLIHY SVGEGT TAKGEVPDGYNVSRLLKQNFLLGLESAAQTSVYFC ASRTDRRYSETQY FGPGTRLLVL

BD[®] OMICS-One Immuno-Oncology Protein Panel

Tumor and adaptive immune system interaction




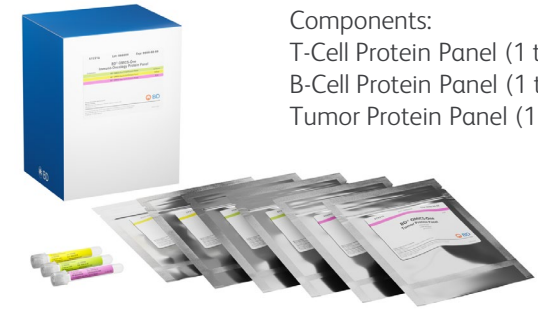
<https://www.mdpi.com/1467-3045/46/1/11>

BD® OMICS-One Immuno-Oncology Protein Panel specificities

Specificity	Clone
CD1d	CD1d42
CD3	UCHT1
CD4*	SK3
CD5	UCHT2
CD8	SK1
CD9	M-L13
CD10	HI10a
CD19	SJ25C1
CD20	2H7
CD21	B-ly4
CD22	HIB22
CD23	EBVCS-5
CD24**	ML5
CD25	2A3
CD26	M-A261
CD27	M-T271
CD28	L293
CD29	MAR4
CD30	BERH8
CD31 (PECAM-1)	WM59
CD34**	581
CD38	HIT2
CD40	5C3
CD43*	1G10
CD44*/**	L178
CD45*	HI30
CD45RA	HI100
CD45RO	UCHL1
CD47	B6H12

Specificity	Clone
CD54	HA58
CD58	1C3
CD62L	DREG-56
CD66	B1.1/CD66
CD69	FN50
CD73	AD2
CD79b	CB3-1
CD80	L307.4
CD90	5E10
CD95**	DX2
CD103	Ber-ACT8
CD117	YB5.B8
CD126	M5
CD127	HIL-7R-M21
CD133	W6B3C1
CD134 (OX40)	ACT35
CD137	4B4-1
CD138	MI15
CD146	P1H12
CD154	TRAP1
CD155	TX24
CD161 (KLRB1)	HP-3G10
CD183	1C6/CXCR3
CD184 (CXCR4)	12G5
CD185 (CXCR5)	RF8B2
CD194 (CCR4)	1G1
CD196 (CCR6)	11A9
CD197 (CCR7)	2-L1-A
CD223 (LAG-3)	T47-530

Specificity	Clone
CD227 (MUC1)	HMFG2
CD268 (BAFF-R)	11C1
CD272	J168-540
CD274 (PD-L1)	MIH1
CD275	2D3/B7-H2
CD278	DX29
CD279 (PD-1)**	EH12.1
CD324 (E-Cad)	67A4
CD325 (N-Cad)	8C11
CD326 (EpCAM)	EBA-1
CD357 (GITR)	V27-580
CD366 (TIM-3)	7D3
c-MET	3D6
EGFR	EGFR.1
EphB2	2H9
HLA-A,B,C*	G46-2.6
HLA-DR* 	G46-6
IgD	IA6-2
IgG	G18-145
IgM	G20-127
Integrin β7	FIB504
Notch1	MHN1-519
Podoplanin	LpMab-17
TCR Vα24-Jα18	6B11
TCRγ/δ	11F2
TIGIT	TgMab-2
Vista	MIH65.rMAb



Components:
T-Cell Protein Panel (1 test/pouch, 2 pouches)
B-Cell Protein Panel (1 test/pouch, 2 pouches)
Tumor Protein Panel (1 test/pouch, 2 pouches)

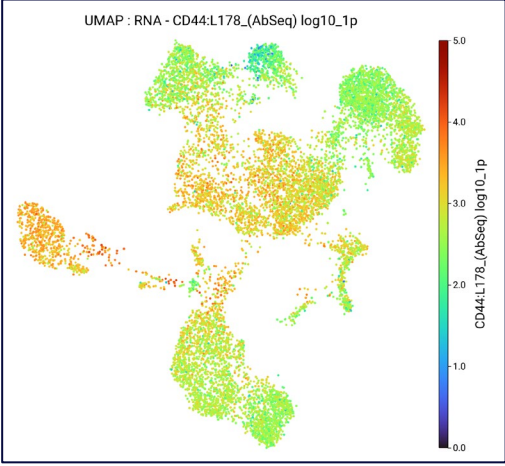
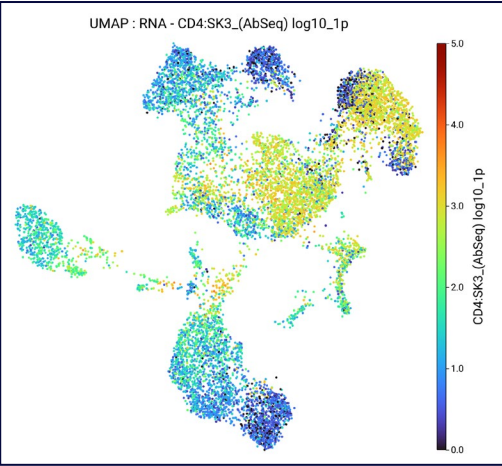
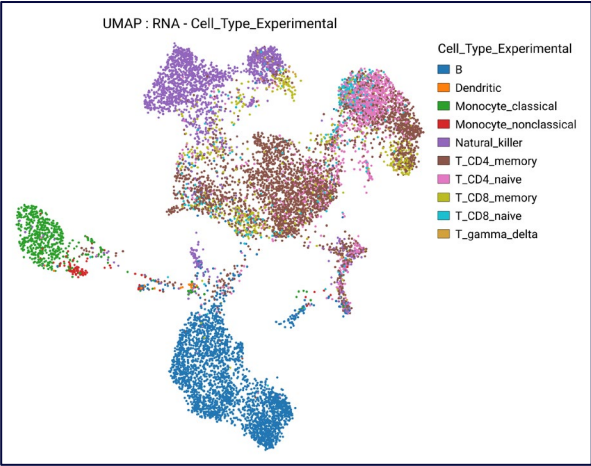
Fewer sequencing reads consumed by CD4, CD43, CD44, CD45, HLA-DR and HLA-A,B,C and more reads allocated to lowly expressed markers

Percent of Total Sequencing Reads Consumed		
Markers	Without SMART panel design	With SMART panel design
Reduction of sequencing reads allocated to primary markers ▼		
CD44	27.14	8.80
HLA-DR	13.42	8.74
CD43	6.07	3.37
HLA-A,B,C	5.61	4.45
CD4	2.10	1.47
CD45	3.19	1.48
Read re-allocation to selected lowly expressed markers ▲		
CD34	3.23	3.40
CD54	2.38	4.33
CD69	3.37	6.90
CD5	2.04	3.25
CD20	0.46	1.21
CD27	1.44	2.07
CD19	0.43	0.75
CD25	1.32	2.72

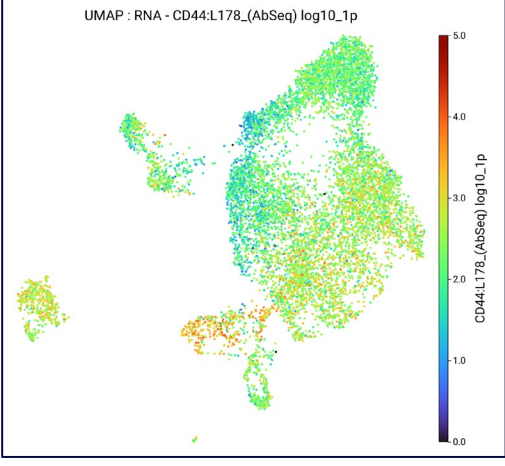
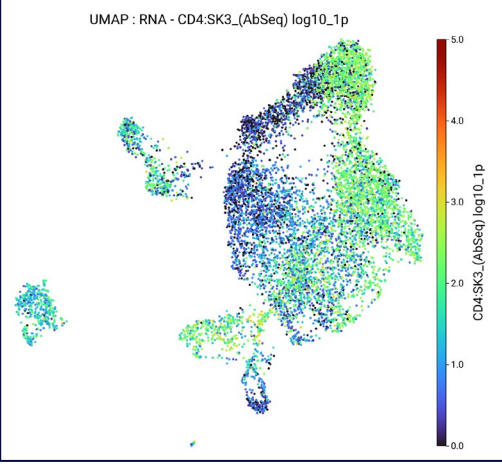
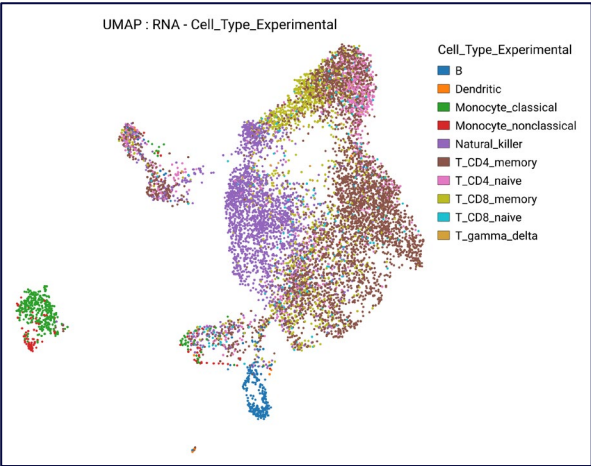
Percent of Total Sequencing Reads Consumed		
Markers	Without SMART panel design	With SMART panel design
CD279	1.16	1.76
CD79b	0.98	1.45
CD196	0.94	1.51
CD95	0.91	1.30
CD73	0.83	1.28
CD3	0.81	1.54
CD38	0.76	1.23
CD194	0.69	1.37
CD366	0.64	1.23
CD9	0.62	1.00
CD47	0.62	1.07
CD155	0.62	0.88
CD23	0.61	0.96
CD272	0.61	1.21
IgM	0.61	0.99
CD275	0.60	1.42

CD4, CD43, CD44, CD45, HLA-DR and HLA-A,B,C detection is not compromised

Without SMART
panel design

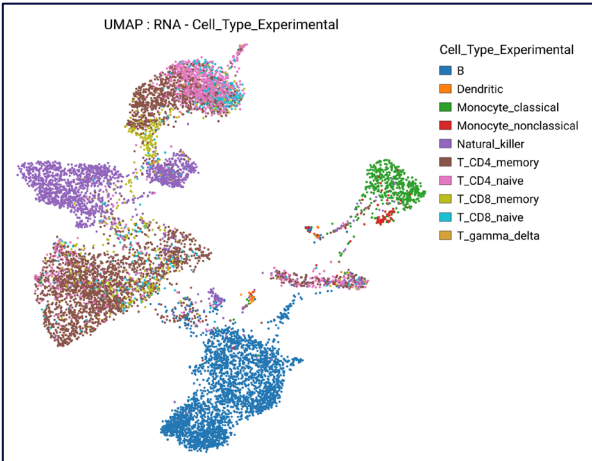


With SMART
panel design

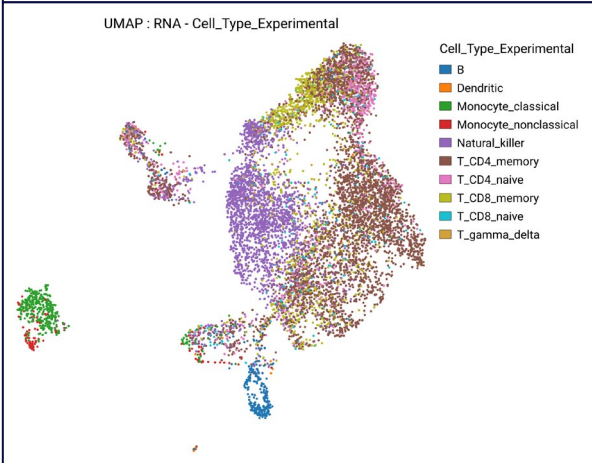


CD4, CD43, CD44, CD45, HLA-DR and HLA-A,B,C detection is not compromised

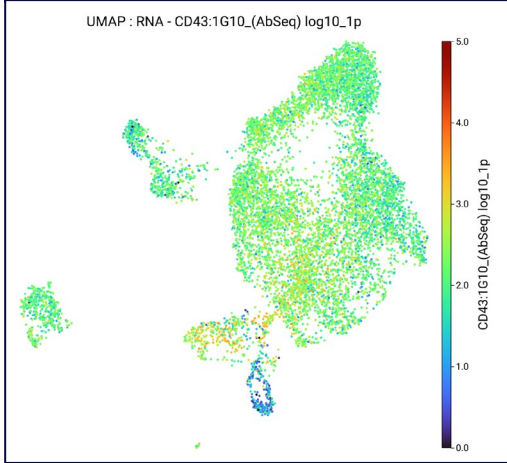
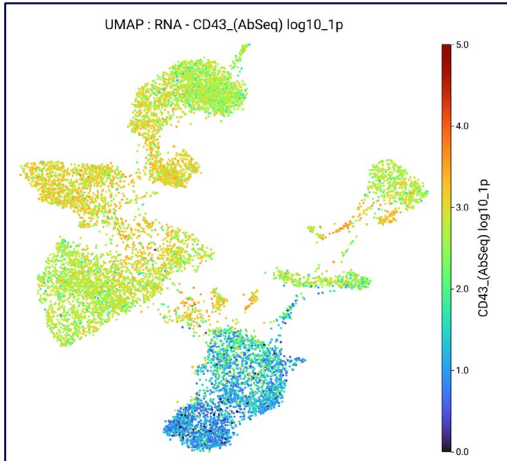
Without SMART
panel design



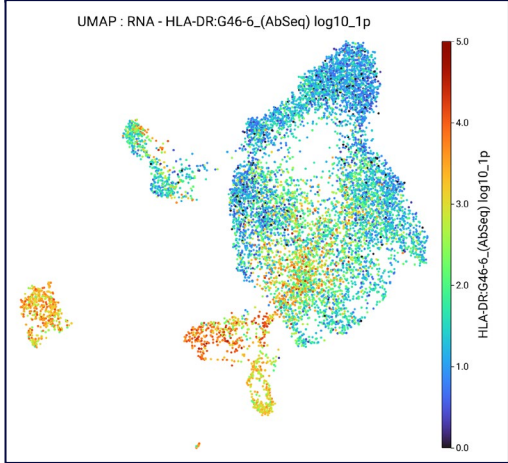
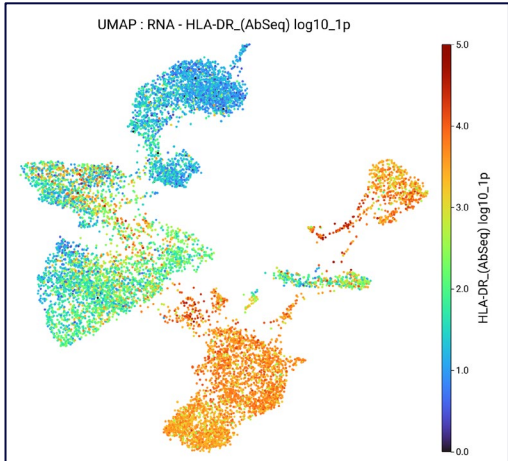
With SMART
panel design



CD43

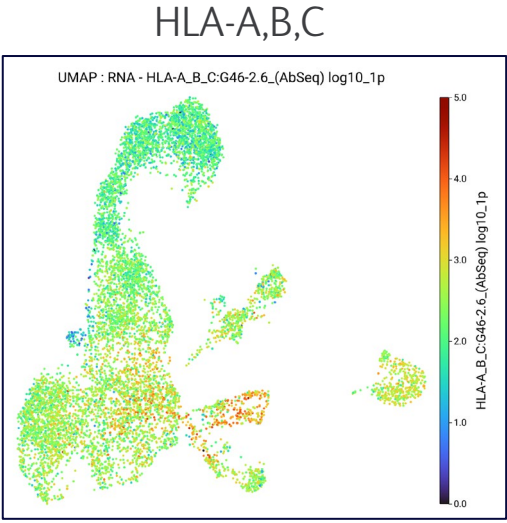
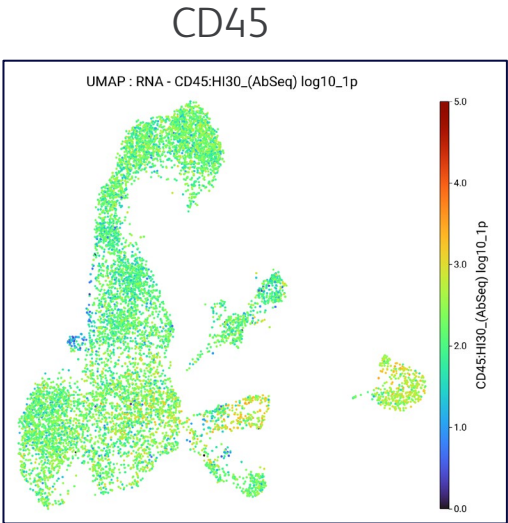
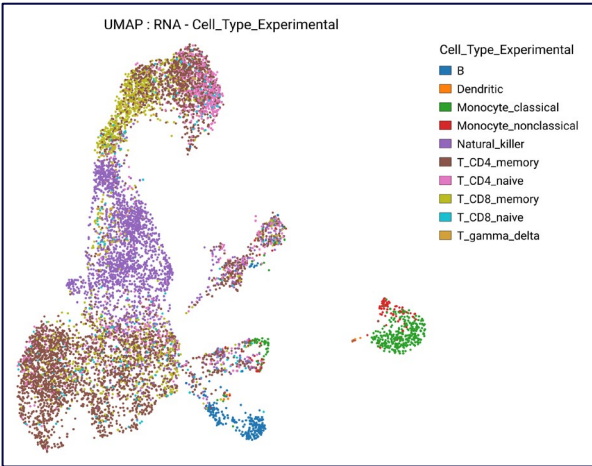


HLA-DR

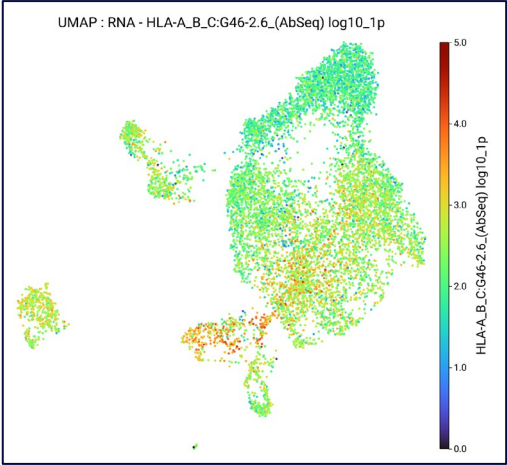
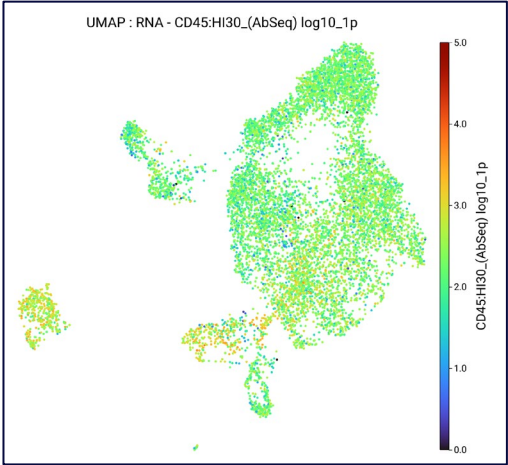
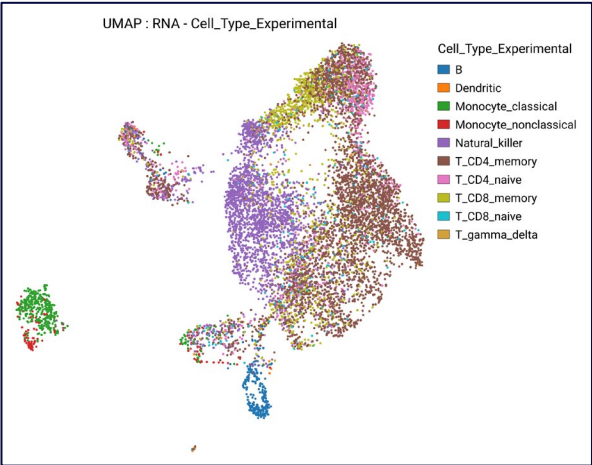


CD4, CD43, CD44, CD45, HLA-DR and HLA-A,B,C detection is not compromised

Without SMART
panel design

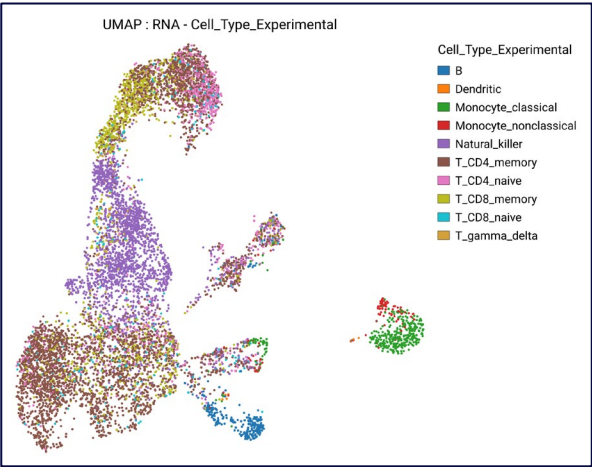


With SMART
panel design

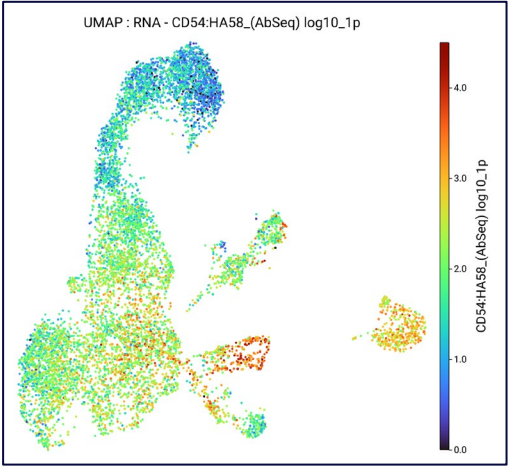


Better resolution of low expressors is found with SMART panel design

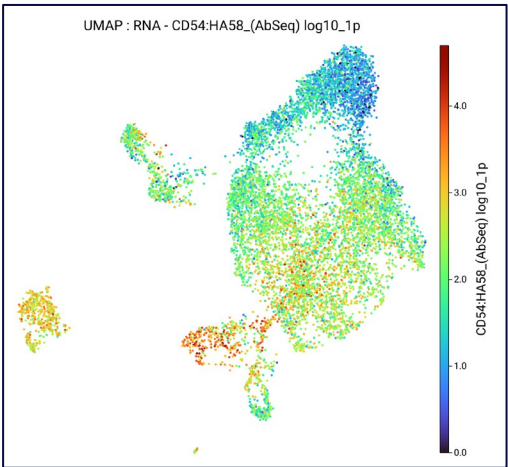
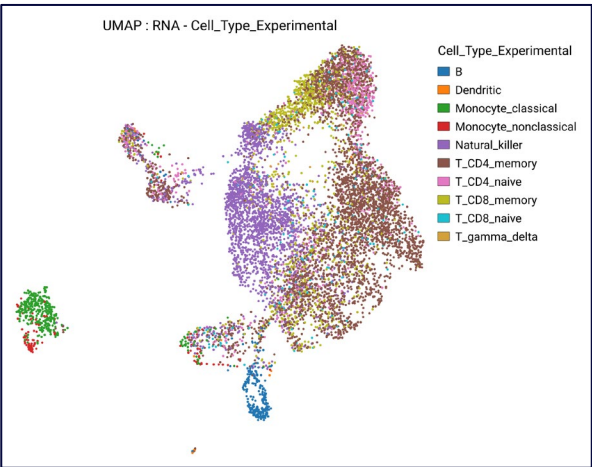
Without SMART panel design



CD54



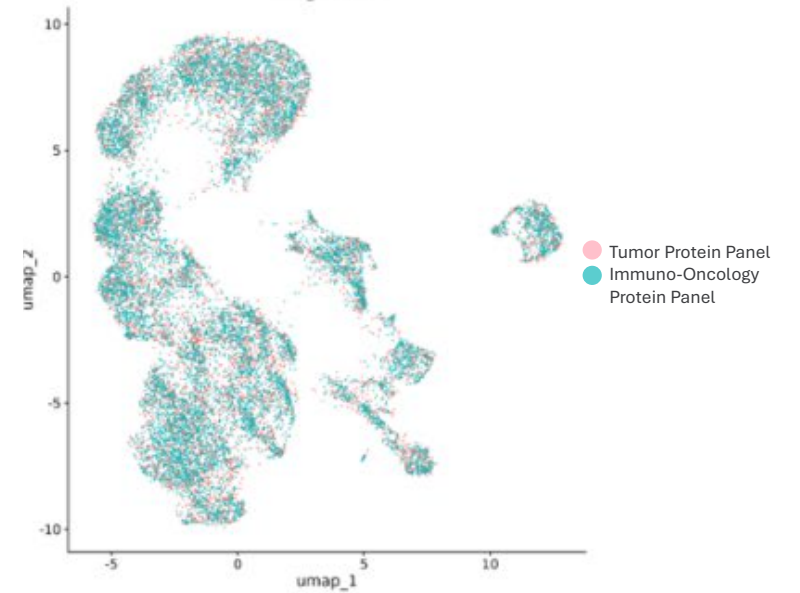
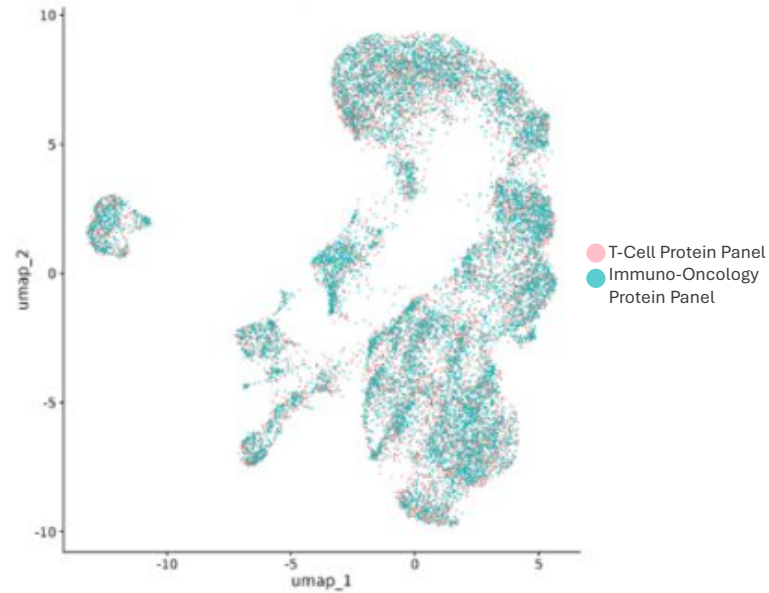
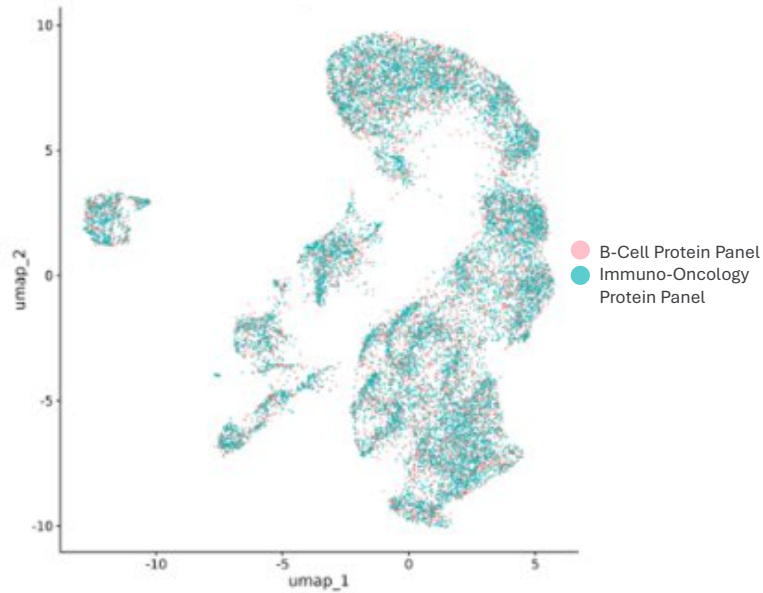
With SMART panel design



Scalable high-plex protein profiling solution with modular panel design

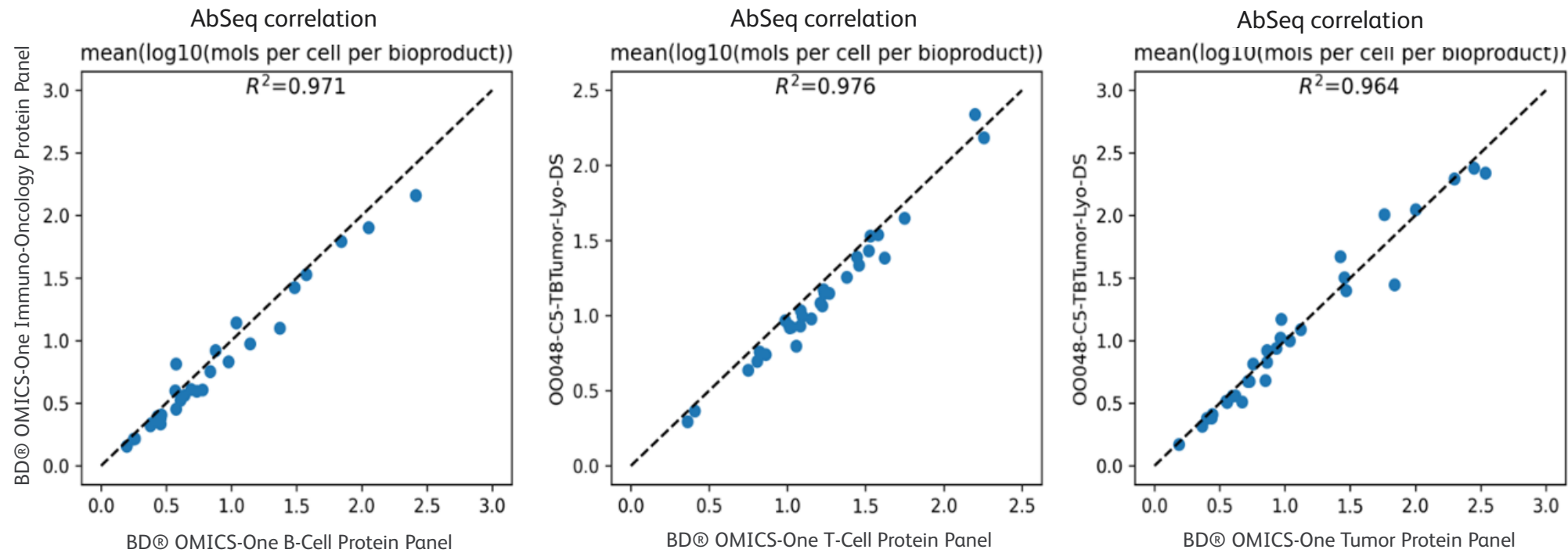
A.

Similar mRNA detection performance between the lyophilized BD[®] OMICS-One Immuno-Oncology Protein Panel versus each lyophilized component panel.

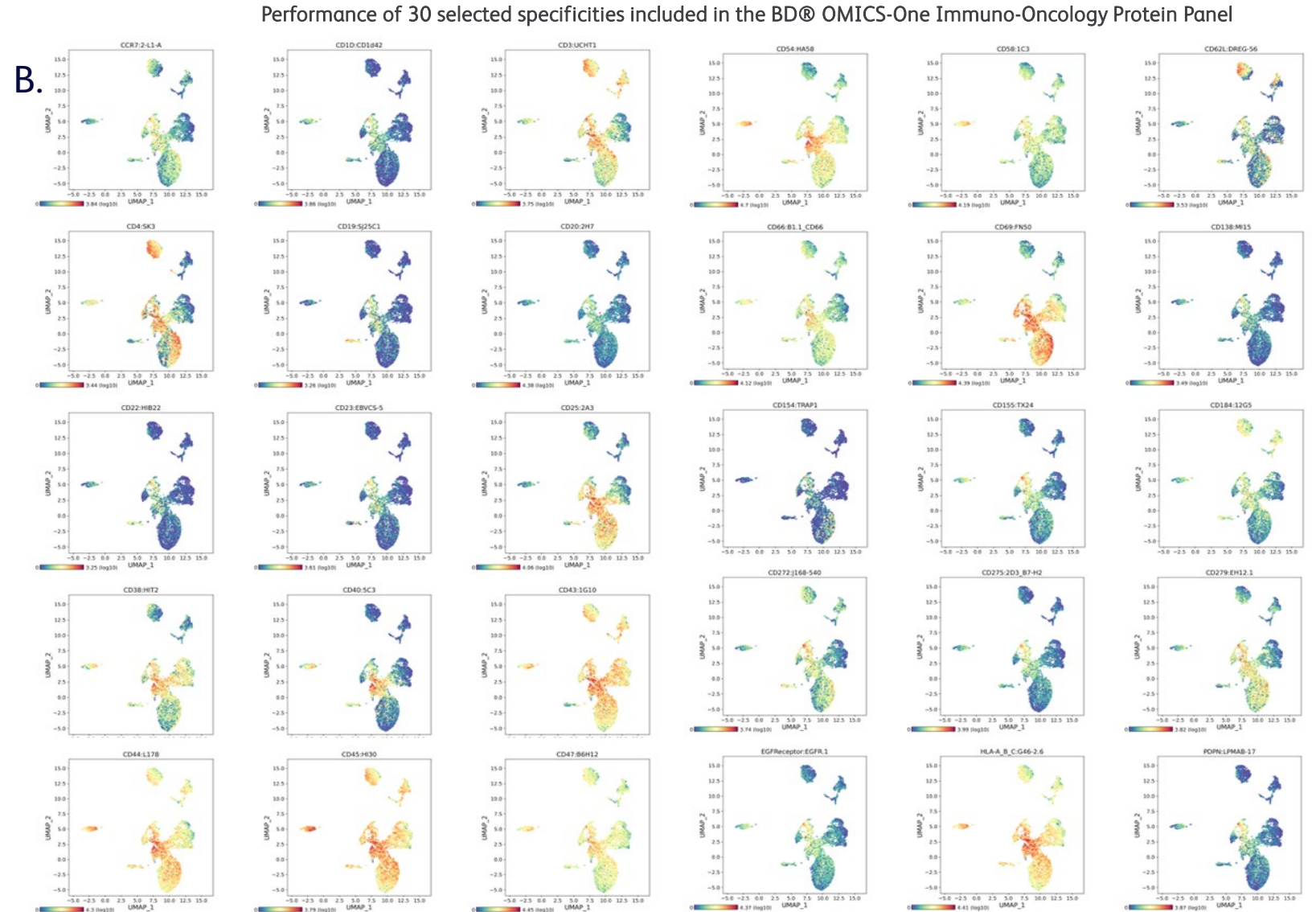
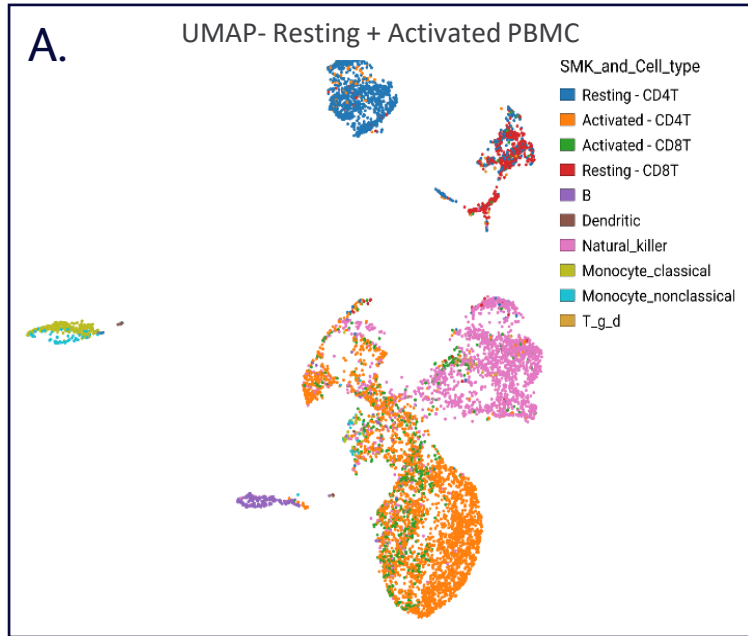


Scalable high-plex protein profiling solution with modular panel design

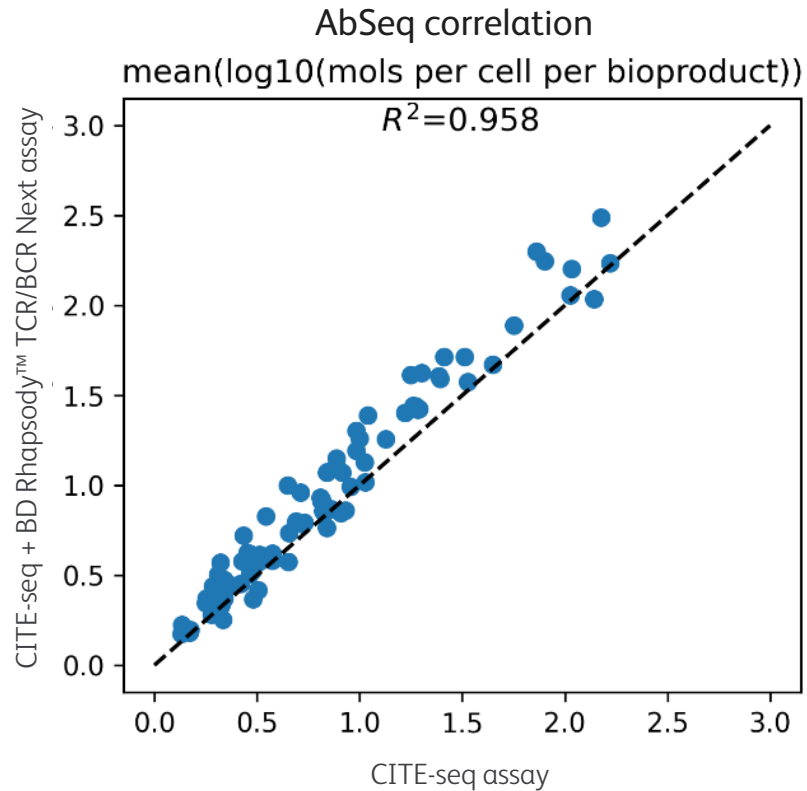
B. Similar protein detection performance between the lyophilized BD[®] OMICS-One Immuno-Oncology Protein Panel versus each lyophilized component panel.



Detect 85 critical adaptive immune and tumor markers in your samples with confidence

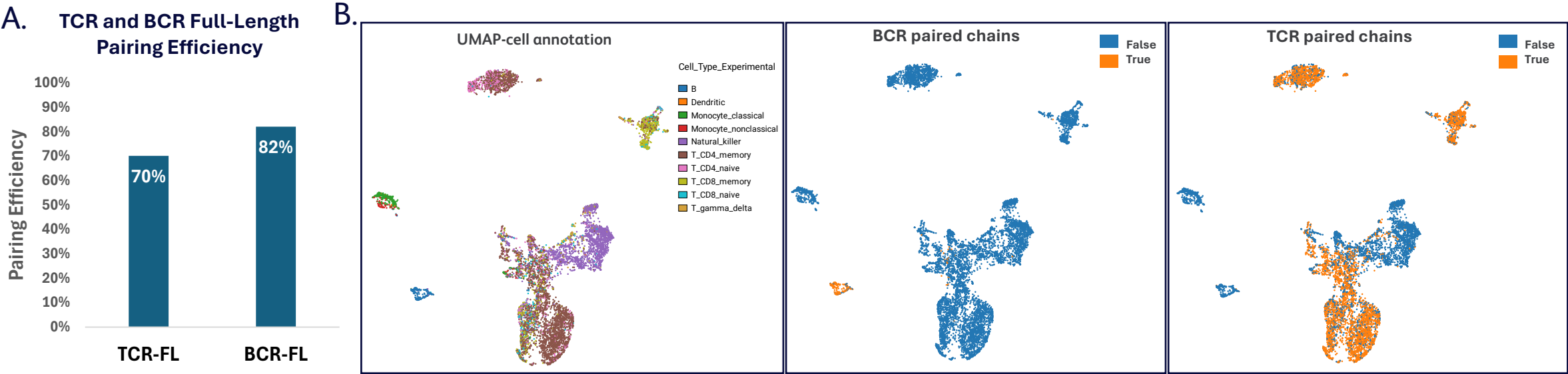


Multiomics enabled: BD[®] OMICS-One Immuno-Oncology Protein Panel is designed to work with WTA and TCR/BCR assays



The addition of BD Rhapsody™ TCR/BCR Next Assay does not impact BD[®] OMICS-One Immuno-Oncology Protein Panel performance

Multiomics enabled: BD[®] OMICS-One Immuno-Oncology Protein Panel is designed to work with WTA and TCR/BCR assays



C. ● FWR1 ● CDR1 ● FWR2 ● CDR2 ● FWR3 ● CDR3 ● FWR4

Cell Index	TCR/BCR Chain	VDJ Translation Trimmed
2937	IG_Heavy	EVQLVESGGGLVKPGGSLRLSCAAS GFAFSTYN MNWVRQAPGKGLEWVSS IGSSTSYT YYAESVKGRFTISRDNAKNSLYLQMTSLRAEDTAVYYC ARDLTNWN DGISGDY WGQGT VTVVSS
2937	IG_Kappa	DIQMTQSPSSLSASLGDRVITITCRAS QSI SRLF LNWYQKPGKAPKLLIY SAS NLQSGVPSRFSFGSGSGTDFTLTISNLQPEDFATYYC QKSY SFPH T FGQGT KLEIK
13092	TCR_Alpha	TQLLEQSPQLSIQEGENLTVYCNS S VFSS LQWYRQEPGEGPVLLVT VVT GGEV KKLKRLTFQFGDARKDSSLHITAAQPGDTGLYC AGAAGT SYGKLT FGQGT ILTVHP
13092	TCR_Beta	DGGITQSPKYLFRKEGQNVTLSC EQN LNHDA MYWYRQDPGQGLRLIYY SQIVND FQKGDIAEGYSVSREKKESFPLTVTSAQKNPTAFYLC ASTRRG VT EAF FGQGT RLTVV
43771	TCR_Alpha	QQPVQSPQAVILREGEDAVINCSS KALYS VHWRQKHGEAPVFLMI LLKGGEQ KGHEKISASFNEKKQSSLYLTASQLSYSGTYFC GTANTN AGKST FGDGT TTLVKP
43771	TCR_Beta	WLGWVVTSSHQVAQMGEVILRCVPI SNHLY FYWYRQILGQKVEFLVS FYNNEI SEKSEIFDDQFSVERPDGSNFTLKIRSTKLEDSAMYFC ASRS AVYNSPLH FGNGT RLTVT
55572	TCR_Alpha	DAKTTQPNMESNEEPPVHLPCNHS TISGTDY IHWYRQLPSQGPEYVIH GLTSN VNNRMASLAIAEDRKSSTLILHRATLRDAVYYC ILRGI QGAQKLV FGQGT RLTINP
55572	TCR_Beta	EAGVAQSPRYKIIERQSVAFWCNPI SGHAT LYWYQQILGQGPKLLIQ FQNGV VDDSQLPKDRFSAERLKGVDSTLKIQALEDASVYLC ASSL AEYGNNEQF FGPGT RLTVL
72959	TCR_Alpha	DAKTTQPTSMDCAEGRAANLPCNHS TISGNEY VYWYRQIHSQGPQYIIH GLKNN ETNEMASLIITEDRKSSTLILPHATLRDTAVYYC IVSYE KL FGTGT RLTIIP
72959	TCR_Beta	GAGVSQTPSNKVTEKGDVLRCDPI SGHTA LYWYRQSLGQGLEFLIY FQNSA PDKSGLPSDRFSAERTGGSVSTLTQRTQQEDSAVYLC ASSL GQGSFNEQY FGPGT RLTVT
79323	TCR_Alpha	GENVEQHPSTLSVQEGDSAVIKCTYS DSASNY FPWYKQELGKRPLIID IRSNVG E KKDQRIAVTLNKTAKHFSLHITETQPEDSAVYFC AASY FGNEKLT FGTGT RLTIIP
79323	TCR_Beta	DTGVSQDPRHKITKRGQNVTFRCMPI SEHNR LYWYRQTLGQGPFLTY FQNEAQ LEKSRLSDRFSAERPFGSFSTLEIQRTEQGDASMYLC ASSA QGGNEQF FGPGT RLTVL

Thank you



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Ordering information

Cat. no.	Product description	Config (Size)	List price (USD)	Shelf life
572178	BD® OMICS-One T-Cell Protein Panel	2 tests/kit	\$600	3 years
572179	BD® OMICS-One B-Cell Protein Panel	2 tests/kit	\$600	3 years
572241	BD® OMICS-One Adaptive Protein Panel	2 tests/kit	\$1,100	3 years
572310	BD® OMICS-One Tumor Protein Panel	2 tests/kit	\$600	3 years
572316	BD® OMICS-One Immuno-Oncology Protein Panel	2 tests/kit	\$1,600	3 years



Required and recommended companion products

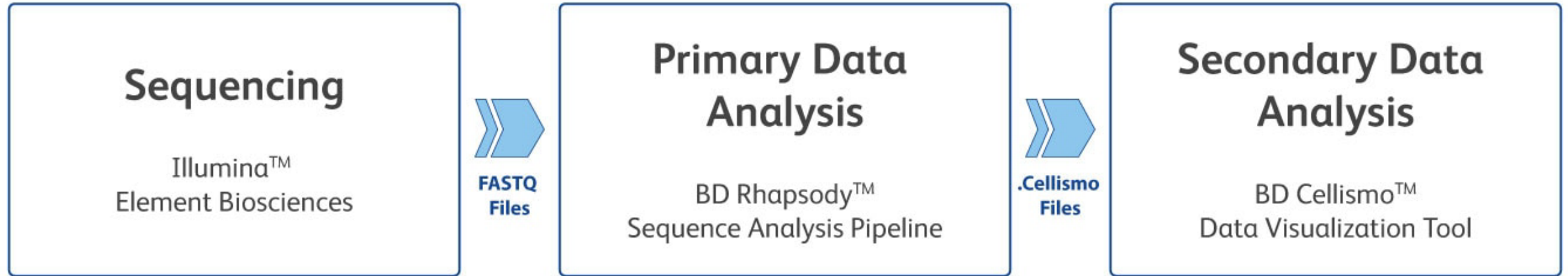
Required companion products

Cat. No.	Product description
554656	BD Pharmingen™ Stain Buffer (FBS)
633801	BD Rhapsody™ Whole Transcriptome Analysis (WTA) Amplification Kit
633773	BD Rhapsody™ cDNA Kit
666262	BD Rhapsody™ 8-lane cartridge
667052	BD Rhapsody™ Enhanced Cartridge Reagent V3
666625	BD Rhapsody™ HT Xpress Package

Recommended companion products

Cat. No.	Product description
667058	BD Rhapsody™ TCR/BCR Next Amplification Kit
633774	BD Rhapsody™ Targeted mRNA and AbSeq Amplification Kit
564220	BD Pharmingen™ Human BD Fc Block
633781	BD® Human Single-Cell Multiplexing Kit
633849	BD® Flex Single-Cell Multiplexing Kit A, Flex Sample Tag 1-6
633850	BD® Flex Single-Cell Multiplexing Kit A, Flex Sample Tag 7-12
633851	BD® Flex Single-Cell Multiplexing Kit A, Flex Sample Tag 13-18
633852	BD® Flex Single-Cell Multiplexing Kit A, Flex Sample Tag 19-24
633701	BD Rhapsody™ Scanner
570742	BD Rhapsody™ Intracellular AbSeq Buffer Kit
570911	BD® OMICS-Guard Sample Preservation Buffer
570750	BD® AbSeq Enhancer
570751	BD® RNase Inhibitor

Sequencing and data analysis



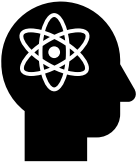
- Recommended sequencing depth for BD® OMICS-One Protein Panels: 300 read pairs/specificity per cell
- Read length: Minimum of 51 × 71 bp

- Get access to the BD Rhapsody™ Sequence Analysis Pipeline on the [Seven Bridges Genomics Platform](#) or on a local installation
- Acquire the AbSeq reference file (.fasta) from the [BD AbSeq Panel Generator](#)
- Set up analysis following the *BD Rhapsody™ Sequence Analysis Pipeline User's Guide* (Doc ID: 23-24580)

- Navigate to the [BD Cellismo™ Data Visualization Tool landing page](#)
- Provide your information*
- Select your operating system
- Install the BD Cellismo™ Data Visualization Tool and agree to the end-user license agreement (EULA)

*Will be used to notify you of updates and bug fixes

Supporting you with your single-cell experiments



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Visit us at scomix.bd.com to view our resource library, learning center and FAQs



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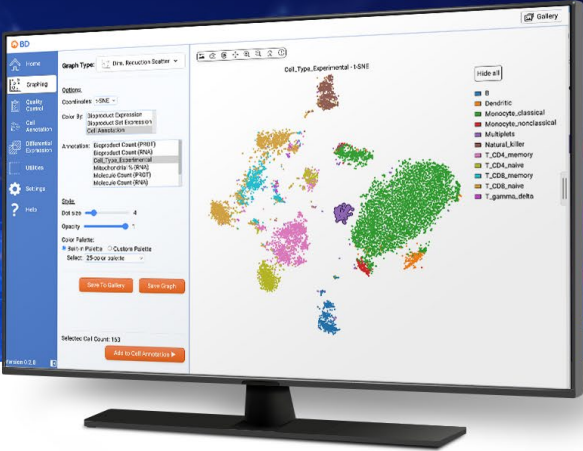
Multiomics



Million-Cell
Throughput



Validated Multiomic
Kits and Protocols



Simple and Free
Bioinformatics