

PROTEOMICS

Supporting Information

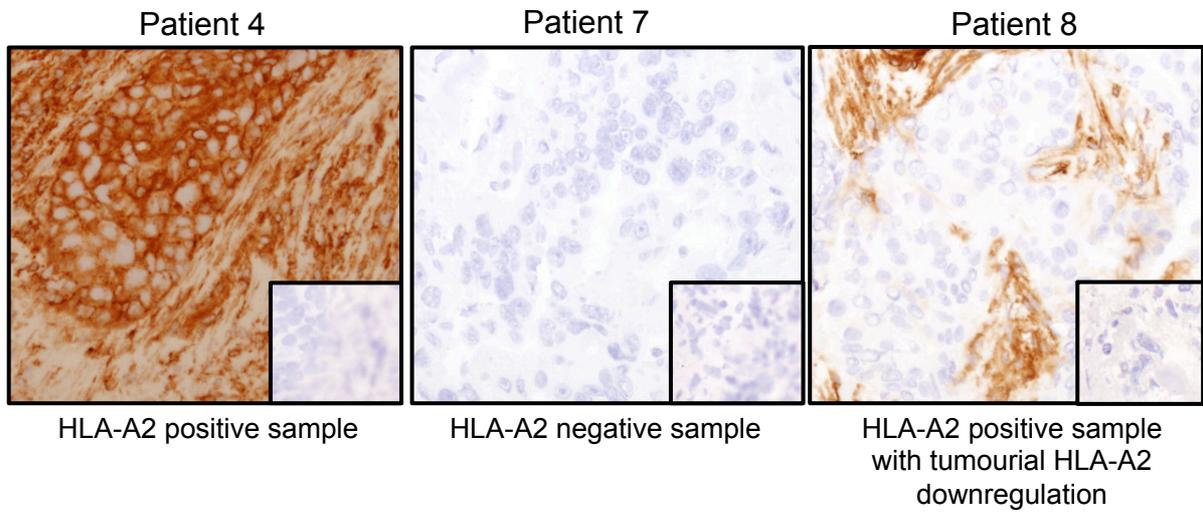
for Proteomics

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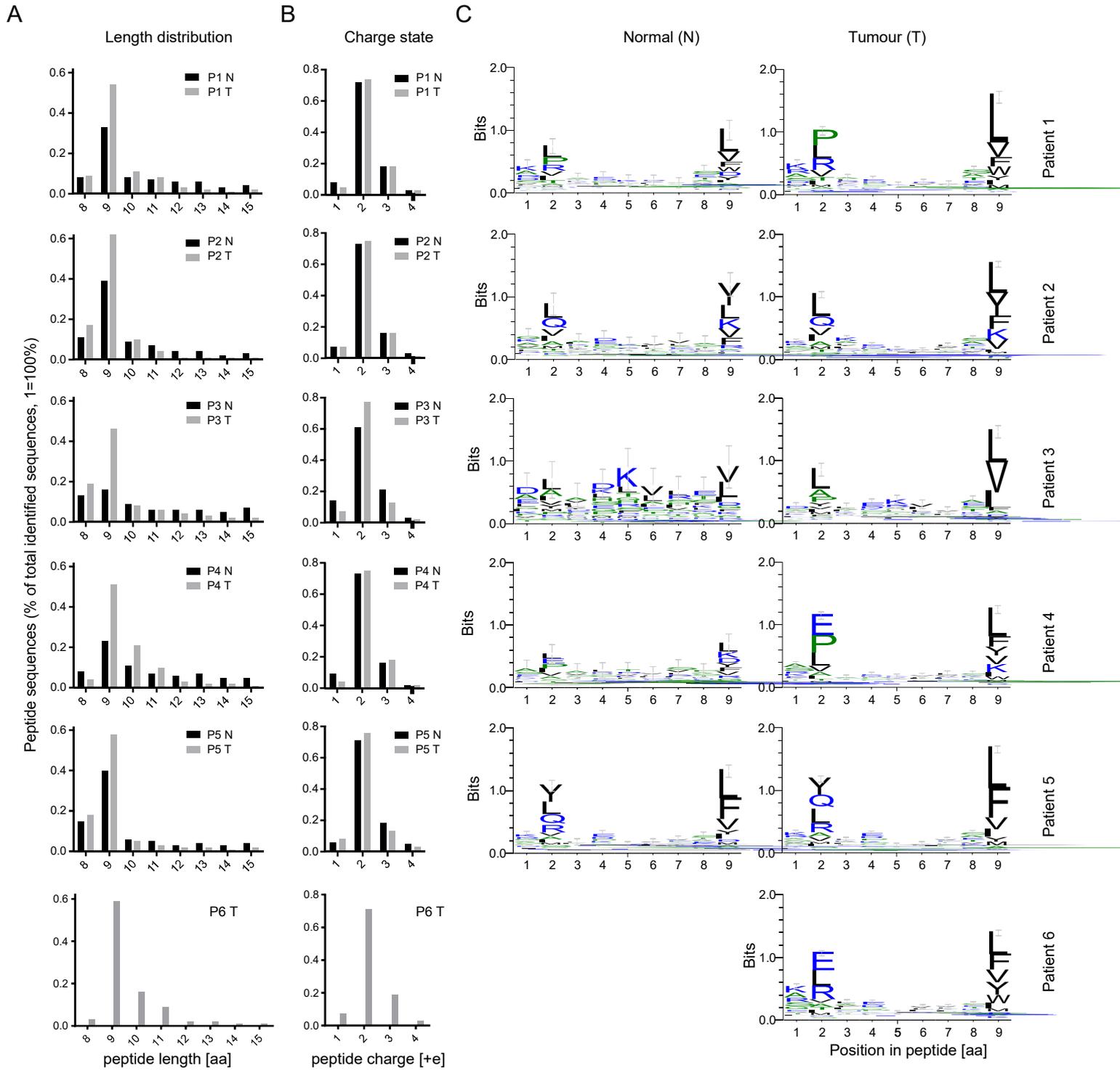
Nicola Ternette, Marloes J. M. Olde Nordkamp, Julius Müller,
Amanda P. Anderson, Annalisa Nicastrì, Adrian V. S. Hill, Benedikt M. Kessler
and Demin Li

**Immuno-peptidomic Profiling of HLA-A2-Positive Triple Negative Breast Cancer
Identifies Potential Immunotherapy Target Antigens**

Supporting Information Figure S1

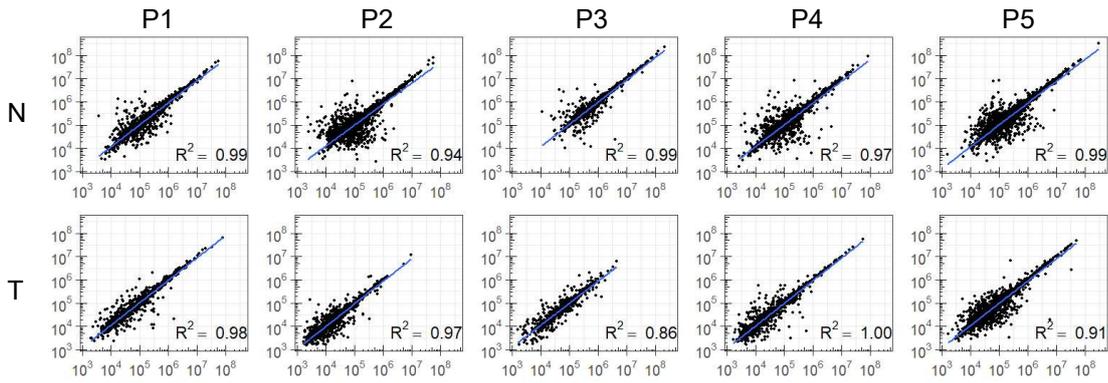


Supporting Information Figure S2

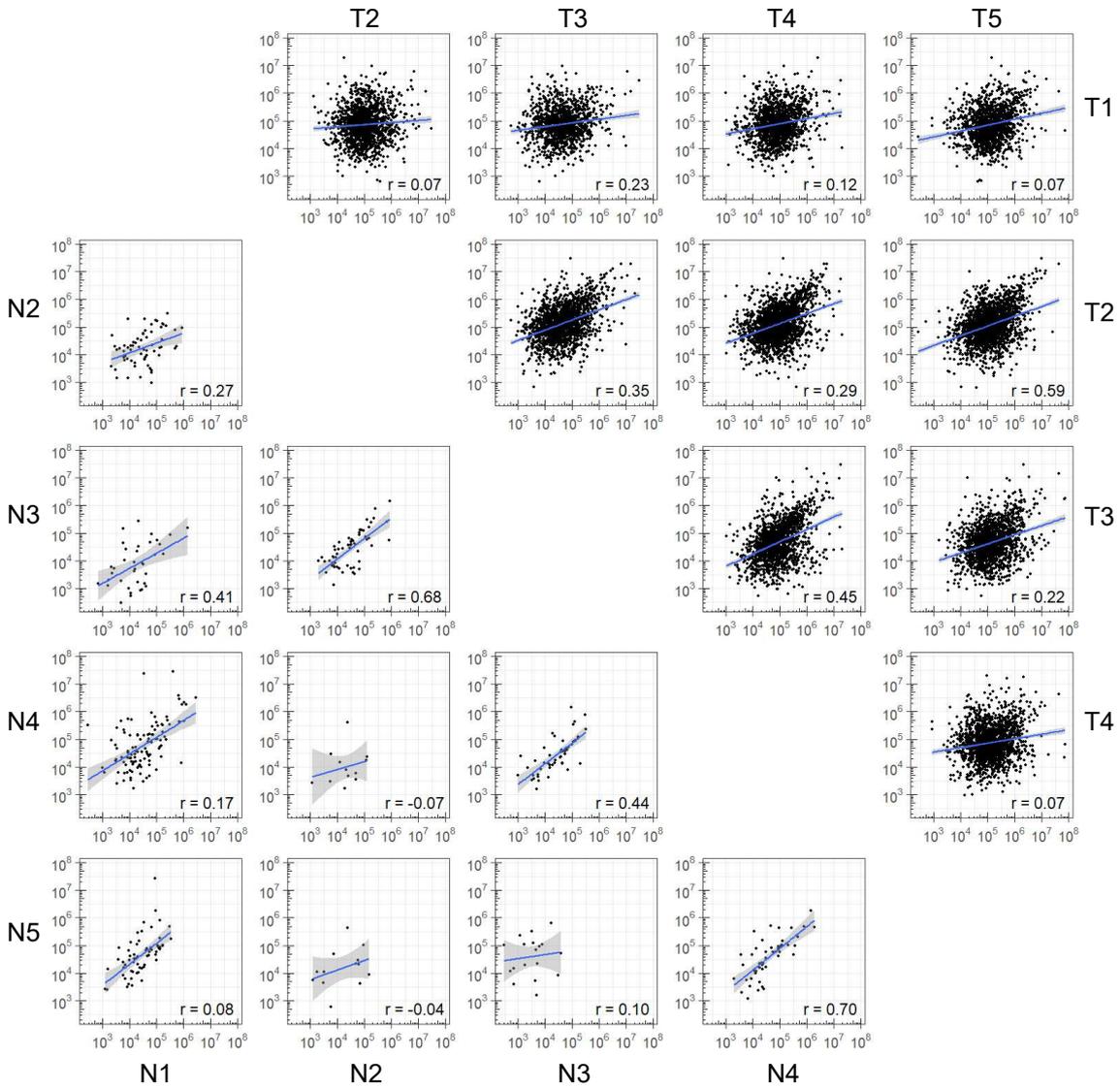


Supporting Information Figure S3

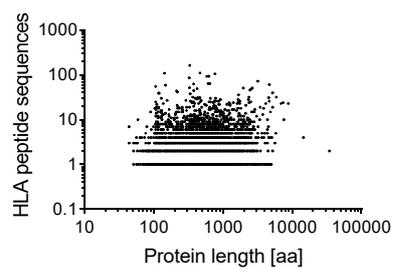
A



B



Supporting Information Figure S5



Supporting Information Figure S6

A

CFL-1

1 MASGVAVSDG VIKVFNDRV RKSSTPEVK KRKKAVLFL SEDKNTLE EKKELIVGDV GQTVDDPYAT FVRLPDKDC
 81 RYALYDQTYE TRSKKEDLV FIFWAPESAP LKSKMIYASS KOATKIKLTLG IKHQLQANCY EEVKQRCITLA EKLGGSAVIS
 161 LEGKPL

IL-32

1 MCFPKVLSD D MKKIKARM M LLTSAQGLG AVNSACDTE TVHGLGPRD KDPALWCQLC L SQRQATER P YDKMNAES
 81 GRQVMSSIA ELEDDFKESQ LETVAAYEE QHPELTPLE KERDGLCRG NRSFVDFVD PATERPESEF CI KVMRHFQA
 161 M LQRLQTHW GVLAVKERV VALVHVAQAL WKQFQSPCCS LSELPMSSFO SYGAPRODKE EL TPKQCEP QSSK

PCNA

1 MFEARLVGGS ILKKVLEAL K DLINACWDI SSSGVLSQM DSSVSLVQL TLRSEGFDY RCDNLANGV NLTMSKILK
 81 CAGNEDIITL RAENADITLA LVFEAPNQEK VSDYEMKLD LOVEQLGIPE QEYSVVMKQ SGEFARICRD LSHIGDAVVI
 161 SCARDGVKFS ASGELGNNGI KLSQTSNVDK EEEAVTIEMN EPVOLTFALR YLNFFTKA TP LSSTVILMS ADVPLVYEK
 241 IADMGLKYY LAKPIEDEEG S

Syntenin-1

1 MSLYPSLEDL KVKVIGQCT AF SANPANPA ILSEASAPIP HDGNLYRLY PELSOY SLS LNNEEIRANV AVSGAPLQG
 81 QLVARPSSIN YMVAPVTGND VGIRRAEIKQ GIREVILCKD QDGKIGLRK SIDNGIFVQL VQANSFASLV GLRFGDQVIQ
 161 INGENCAGS SDKARKVLKQ AF SEKITMTI R RFFERITIT M HKDSTGHV G FIKNGKITS IVKDSAA RN GLTNEICE
 241 INQNVIGLK DSQIADILST SGTVVTITM PAFIFEHIIK RMAPSINM SL MOHTIPEV

RPN-2

1 MAPPGSSIVP LLALTIASL MALTPTHYLT KHDVERLKAS LDRPPTNLES AFYSIVGLSS LGAQVPDAMK ACTYIRSNLD
 81 PSNVDSLVA AQASQALSGC EISISNETKD LLLAAVSEDS SVTQIYHAVA ALSGGLEIA SQFALSALTA RLKEETVLA
 161 TVQALQTASH LSQQADLRSI VEEIEDLVAR LDELGGVYLQ FEEGLETTAL FVAATYKMD HVGTPEPSINE DQVIQMNAI
 241 FSKKNFELS KAFSVASAAA VLSRNRVYVP VVVVPEGSAS DTHEQAILRL QVTNVLSPQL TQATVKLEHA KSVASRATVL
 321 QKTSF TPVGD VFELNFMNVK FSSGYDPLV EVEGDNRVIA NTVLELVKVIS TEVGINVDL STVKDKQ SIA PKTRVTFPA
 401 KAKGTFIADS HONEALFFQL VDVNTGAEL T PHQTFVRLIN QKTQEVVEV AEPDNKNVY FELDTSERKI EFDASGTYT
 481 LYLIIGDQTL KNPILMNVD VVIKPEEAA PSTVLSQNLF TPKEQIHLF REPEKRPVY VNTFTALLL SPILLFALM
 561 ITRIGANVSNF TPAFSTIIFH LGHAAMGLM YVYVQLNRP QTLKYLALG SVTLA NSM LAQQAVR TA H

B

