



Article Alterations of NK Cell Phenotype in the Disease Course of Multiple Myeloma

Tatiana Pazina, Alexander W. MacFarlane IV, Luca Bernabei, Essel Dulaimi, Rebecca Kotcher, Clinton Yam, Natalie A. Bezman, Michael D. Robbins, Eric A. Ross, Kerry S. Campbell and Adam D. Cohen

Supplementary Material



Figure S1. Representative gating and staining of NK cells. (**A**) Gating. strategy. PBMCs from a MM patient were stained for 20 min on ice. Gates were applied to select single cells, propidium-negative viable cells, lymphocytes and then lymphocyte subsets.NK cell subsets were further divided into CD45⁺CD3⁻CD56^{bright} and CD56^{dim} NK cells. (**B**) Representative staining of CD56^{bright} vs CD56^{dim} NK cells and percentage of cells staining forpositive expression of CD57, CD69, DNAM1, NKG2D, and SLAMF7 on NK cells from blood. Gates for % positive cells were set where <1% of cells were gated in isotype control or.unstained control samples (grey).



Figure S2. Effects of elotuzumab on NK cell degranulation. PBMCs were incubated alone, with elotuzumab (Elo), with MM.1R target cells, or with target cells and Elo for 2 h at 37 °C and stained for percent LAMP-1 (CD107a)⁺ cells. NK cells were gated as CD45⁺CD3⁻CD56^{dim} cells after the viability gate. Horizontal lines mark median values, and statistics were calculated with unpaired Mann-Whitney test.



Figure S3. BM gating. (**A**) BM cells were purified from BM aspirates of ND patient and stained for 20 min on ice. Gates were applied to select single cells, propidium-negative viable cells, MM tumor cells (left box gate) and lymphocytes (right box gate). MM cells were identified as CD45^{dim}, CD38⁺⁺, CD138⁺, CD56^{+/-}, CD19^{+/-}. Lymphocytes subsets were further divided into CD19⁻CD56⁺ NK and CD56⁺ T cells that were used as negative population for GITR-L, ULBPs staining in B. (**B**) Example of SLAMF7, GITR-L, ULBP 1, ULBP 2,5,6, ULBP 3 staining on viable MM cells gated in (**A**). For SLAMF7 staining unstained cells are shown in grey.



Figure S4. Correlation of NKG2D expression on BM NK cells with ULBP3 ligand expression on MM cells in BM of ND patients. Each symbol is an individual patient. Statistics were calculated with Spearman's rank correlation coefficient, lines are linear least squares fit for visual purposes only. Comparisons are of MFI of NKG2D expression on CD56^{bright} (left) or CD56^{dim} (right) blood NK cell populations vs. % MM cells in BM that are ULBP3⁺.



Figure S5. Kaplan-Meier survival plots showing time to progression based on expression of various NK cell markers for the combined ND, RR, and pSCT MM patients divided into tertiles. Patients were divided into tertiles based on their (**A**) % CD56^{bright} and CD56^{dim} NK cells that are DNAM1⁺, (**B**) % CD56^{bright} NK cells that are CD69⁺, and (**C**) % CD56^{dim} NK cells that are CD57⁺ in blood. Dark solid lines represent high expression tertile, dashed dark lines designate medium expression tertile, and solid grey line is low expression tertile for each surface marker. M-C = Mantel-Cox statistical analysis of tertiles, CPH = Cox-Proportional Hazards test performed on all data points. Numbers of patients in each tertile that are newly diagnosed (ND), relapsed/refractory (RR), or post stem cell transplant (pSCT) are indicated adjacent to each plotted line.



Figure S6. Kaplan-Meier survival plots showing time to progression as a function of SLAMF7 expression on blood NK cells in pSCT and RR MM patients. The pSCT (**A**) and RR (**B**) patients were divided into halves above and below median based on the dMFI of SLAMF7 expression on their CD56^{bright} (left) and CD56^{dim} NK cells (right) in peripheral blood. Solid lines designate the upper half and dashed lines are the lower half of SLAMF7 expression. M-C = Mantel-Cox statistical analysis of halves, CPH = Cox-Proportional Hazards test performed on all data points.

Table S1. NK cell antibody staining panel for flow cytometry. Each column represents one staining tube and each row is a fluorescent channel on the flow cytometer. Monoclonal antibodies to markers are listed, followed by clone name and manufacturer. (EB = eBioscience, BL = Biolegend, BD = BD Pharmingen, BC = Beckman Coulter, BMS = Bristol-Myers Squibb).

Fluoro- phore	NK #1	NK #2	NK #3	T/NK	т	lsotype #1	lsotype #2	lsotype #3
FITC	CD19 H1B19 (EB)	CD56 NCAM 16.2 (BD)	KIR2DL1/KIR2DS1 EB6B (BC)	CD56 NCAM 16.2 (BD)	CD25 2A3 (EB)	CD56 NCAM 16.2 (BD)	CD56 NCAM 16.2 (BD)	CD56 NCAM 16.2 (BD)
PE	CD57 NK1 (BD)	KIR3DL1/ KIR3DS1 Z27.3.7 (BC)	KIR2DL1/CD158a 143211 (R&D)	CD4 OKT4 (BL)	CD62L DREG-56 (BL)			
APC	SLAMF7 Elo- tuzumab (BMS)	DNAM-1 11A8 (BL)	KIR2DS4 FES172 (BC)	TIGIT MBSA43 (EB)	PD-1 MIH4 (EB)	hulgG1-κ (South- ern Bio- tech)	mlgG1-к (EB)	
PerCP/ Cy5.5	CD45 2D1 (EB)	CD45 2D1 (EB)	CD45 2D1 (EB)	LAG3 3DS223H (EB)	CD45RA HI100 (BL)	CD45 2D1 (EB)	CD45 2D1 (EB)	CD45 2D1 (EB)
Cy7/APC	CD3 SK7 (BD)	CD3 SK7 (BD)	CD3 SK7 (BD)	CD3 SK7 (BD)	CD3 SK7 (BD)	CD3 SK7 (BD)	CD3 SK7 (BD)	CD3 SK7 (BD)
Alexa- Fluor 700	CD137 4B4-1 (BL)	KIR3DL1 DX9 (BL)	CD11a MEM25 (Exbio)	CD8 HIT8a (BL)	CD8 HIT8a (BL)	CD16 3G8 (BL)	mlgG1-κ (BL)	
QDot 605		NKp44 P44-8 (BL)	KLRG1 2F1/KLRG1 (BL)	TIM3 F38-2E2 (BL)	CD127 eBioRDR5 (EB)		mlgG1-κ (BL)	Syrian Ham. IgG (BL)
Cy7/PE	CD56 NCAM 16.2 (BD)	NKp30 AF29- 4D12 (EB)	CD56 NCAM 16.2 (BD)	OX40 Ber- ACT35 (BL)	CD56 NCAM 16.2 (BD)	NKG2D 1D11 (BL)	mlgG1-κ (BL)	
PerCP	Propidium Iodide	Propidium Iodide	Propidium Iodide	Propidium Iodide	Propidium Iodide	Propid- ium Io- dide	Propid- ium Iodide	Propid- ium Iodide
Pacific Blue	CD69 (FN50) (BL)	GITR eBioAITR (EB)	NKp46 9E2 (EB)	ICOS C398.4A (BL)	CD4 OKT4 (BL)	CD19 HIB19 (BD)	mlgG1-κ (BL)	Arm. Ham. IgG (BL)
Brilliant Violet 510			PD-1 EH12.2H7 (BL)	PD-1 EH12.2H7 (BL)	CD28 CD28.2 (BL)		mlgG1-κ (BL)	

Table S2. Ligands on MM cells antibody staining panel for BM samples assessed by flow cytometry. Each column represents one staining tube and each row is a fluorescent channel on the flow cytometer. Monoclonal antibodies to markers are listed, followed by clone name and manufacturer. (EB = eBioscience, BL = Biolegend, BD = BD Pharmingen, BC = Beckman Coulter, BMS = Bristol-Myers Squibb, RD = R&D Systems).

Fluorophore	#1	#2	#3	#4	
FITC	CD56	HLA-A,B,C	CD56	ULBP1	
	NCAM 16.2	W6/32	NCAM 16.2	FAB1380G	
	(BD)	(BL)	(BD)	(RD)	
PE	CD137-L	Nectin 2 (CD112)	GITR-L	ULBP3	
	5F4	TX3	109101	FAB1517P	
	(BL)	(BL)	(RD)	(RD)	
APC	MICA/MICB	PVR (CD155)	SLAMF7	ULBP2,5,6	
	6D4	300907	Elotuzumab	FAB1298A	
	(BL)	(RD)	(BMS)	(RD)	
PerCP/Cy5.5	CD45	CD45	CD45	CD45	
	2D1	2D1	2D1	2D1	
	(EB)	(EB)	(EB)	(EB)	
Cy7/APC	CD38	CD38	CD38	CD38	
	HIT2	HIT2	HIT2	HIT2	
	(BL)	(BL)	(BL)	(BL)	
AlexaFluor 700	CD19	CD19	CD19	CD19	
	H1B19	H1B19	H1B19	H1B19	
	(EB)	(EB)	(EB)	(EB)	
QDot 605	PDL2 MIH18 (EB)	ICOS-L MIH12 (EB)		BCMA BAF193 (RD)	
Cy7/PE	PDL1 29E2A3 (BL)	CD56 NCAM 16.2 (BD)		CD56 NCAM 16.2 (BD)	
PerCP	Propidium	Propidium	Propidium	Propidium	
	Iodide	Iodide	Iodide	Iodide	
Pacific Blue	CD138	CD138	CD138	CD138	
	MI15	MI15	MI15	MI15	
	(BD)	(BD)	(BD)	(BD)	