July 27, 2022

Cytokine-Based Drug Development Summit, Boston MA

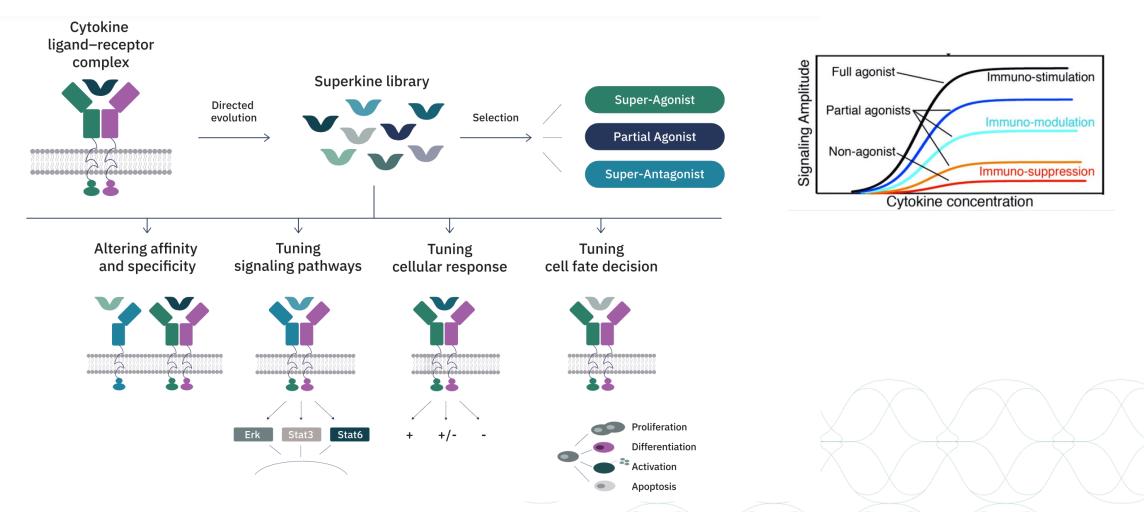
Co-Stimulation of Adaptive and Innate Immune Cells to Achieve Clinical Benefit with MDNA11, a Long-Acting 'Betaonly' IL-2 Super-Agonist

Fahar Merchant, PhD President & CEO Medicenna Therapeutics



Directed Evolution + Yeast Display = Tunable Superkines

> Platform has generated a library of Superkines with diverse immune modulating capabilities



Overview of Superkine Platform

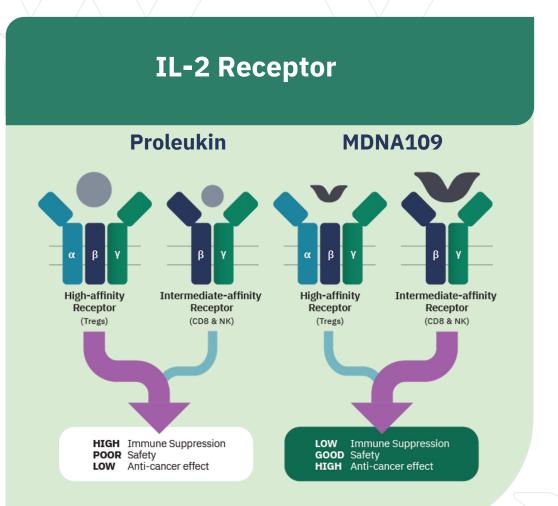
Superkine Platform: Drug Discovery Engine	 Directed evolution enhances the desired properties of IL-2, IL-4, & IL-13 to generate Superkines Protein fusion can improve PK, add an MOA, or confer new capabilities to Superkines IL-2, IL-4, & IL-13 are known to modulate immune activity against 2,000 different diseases
MDNA11: "Beta-only" & Long-acting IL-2 Super- agonist in Phase 1/2	 Super-agonist against IL-2R, a clinically validated anti-cancer target Enhanced IL-2Rβ binding and lack of IL-2Rα affinity position MDNA11 to be best-in-class Preferential stimulation of CD8⁺ T and NK Cells Synergistic with immune checkpoint inhibition Strong memory response with low immunogenicity risk



BiSKIT Platform: Bifunctional SuperKines for ImmunoTherapy

- Fusion of two Superkines or a Superkine and an antibody (e.g. a checkpoint inhibitor)
- Incorporate **two synergistic MOAs** into a single molecule

Targeting IL-2 Receptor Subunits in Cancer Therapy



- > The IL-2 receptor (IL-2R) consists of three subunits
 - CD25 (IL-2Rα)
 - CD122 (IL-2Rβ)
 - CD132 (IL-2Rγ)

Stimulation of CD122

• Key for the activation of cancer-killing immune cells such as CD8+ T cells, naïve T cells, and NK cells.

Stimulation of CD25

- Leads to the activation of immunosuppressive Tregs, which abrogate the anti-tumour response
- Causes extreme toxicity (i.e., pulmonary edema, VLS)

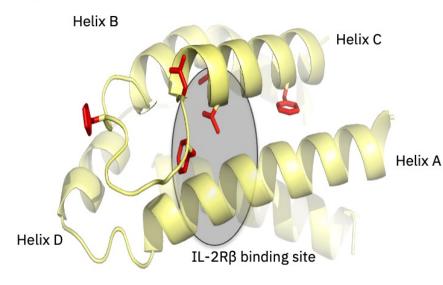
Proleukin® (recombinant human [rh] IL-2), which selectively stimulates CD25, is approved for the treatment of metastatic melanoma and renal cell carcinoma

MDNA109 is a First-Generation IL-2 Super-Agonist

LETTER

nature

Exploiting a natural conformational switch to engineer an interleukin-2 'superkine'

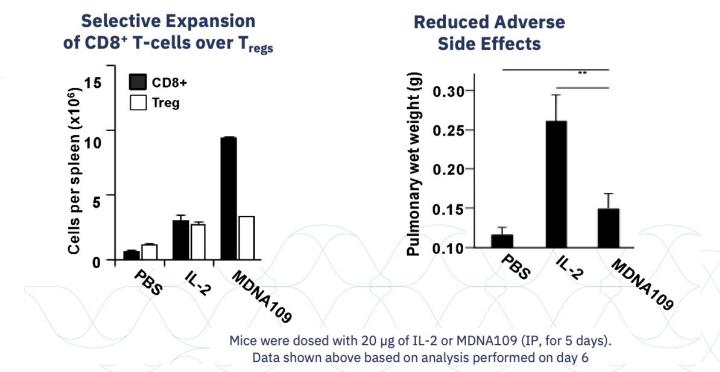


Core Mutations: L80F, R81D, L85V, I86V, I92F

Levin et. al, Nature, 2012 484(7395): 529-533.

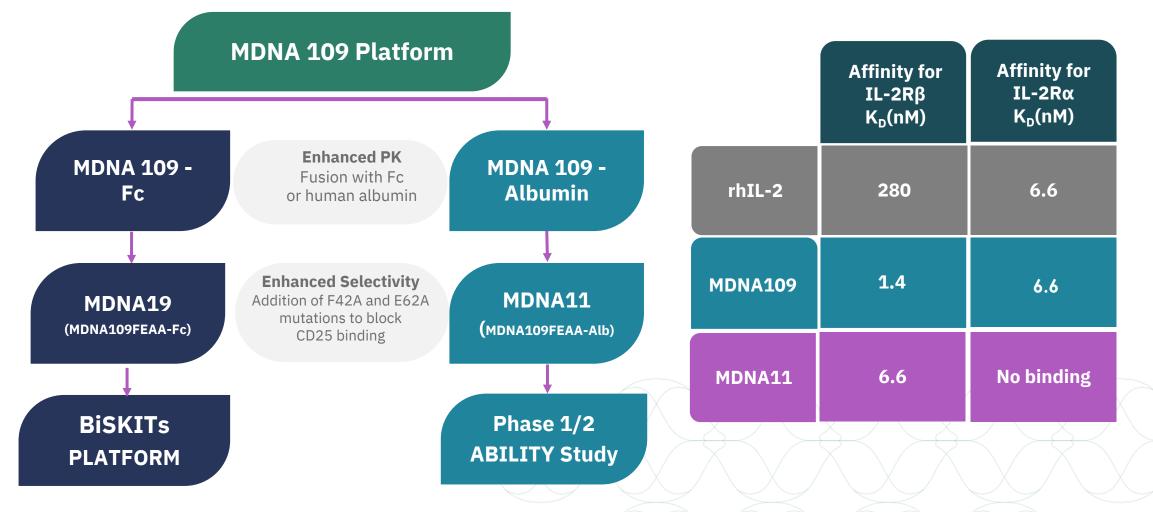
Enhanced Affinity for CD122 (IL-2Rβ); Retains binding to CD25 (IL-2Rα)

SPR data K _D (nM)	CD122 (IL-2Rβ)	CD25 (IL-2Rα)
IL-2	280	6.6
MDNA109	1.4	6.6



MDNA109 Has Been Engineered to Optimize PK & Selectivity

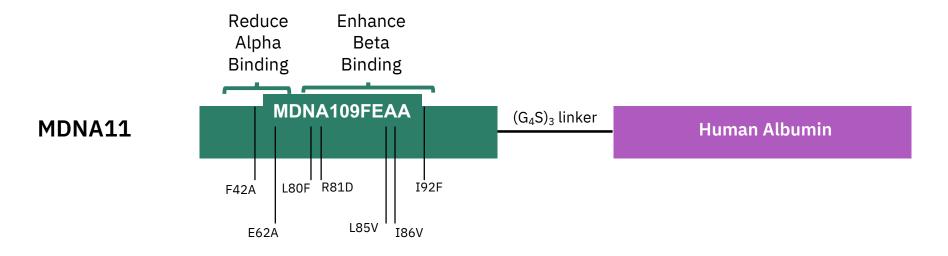
> MDNA11 Has Enhanced IL-2Rβ Affinity, No IL-2Rα Binding, Extended Half-Life and Improved Bioavailability



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Key Differentiators of MDNA11

- > Does not rely on Pegylation or steric hindrance to modulate activity
- Ease of manufacture ensuring consistent batch to batch reproducibility
- Highly versatile backbone aiding design of bespoke immune modulation



Differentiated 'Beta-Only' IL-2 Agonist

- Enhanced affinity for CD122
 - Potentiate CD8 T and NK cells
- No binding to CD25
 - $\,\circ\,\,$ Reduced capacity to stimulate T_{regs}
 - Improved safety profile

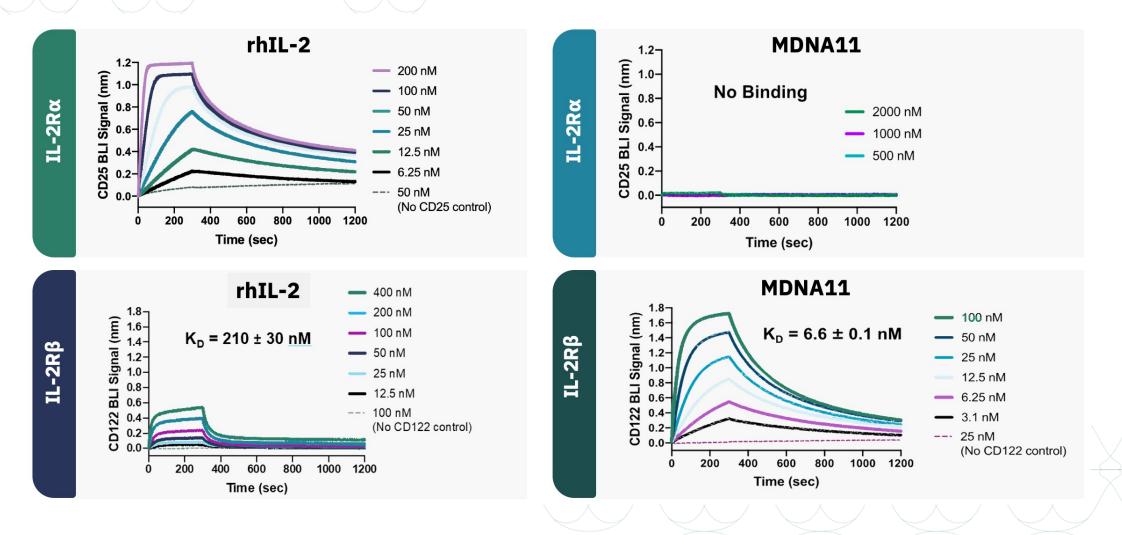
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Fusion to Human Albumin

- Extends in vivo half-life
 - Reduced clearance by kidney filtration
 - Leveraging FcRn recycling
- Potential for accumulation at tumor site and tumor draining lymph nodes
 - Enhanced therapeutic response

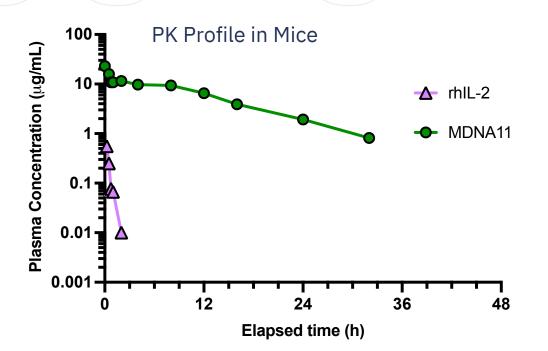
MDNA11 Receptor Binding is Highly Selective Unlike IL-2

> No IL-2Rα (CD25) Binding and Enhanced Affinity and Selectivity for IL-2Rβ (CD122) Compared to rhIL-2



MDNA11 Exhibits Durable Tumor Accumulation In Vivo

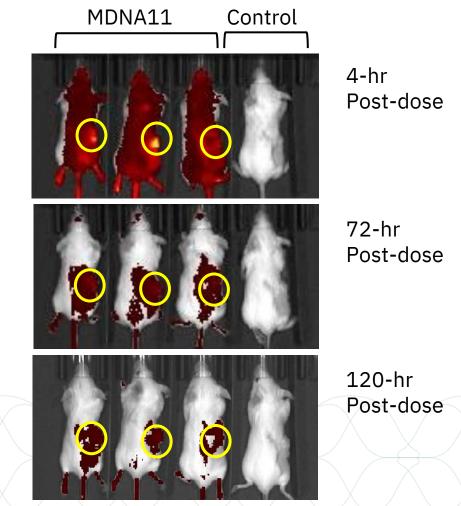
Improves Half-Life by ~25 fold in mice and tumor bioavailability for at least 120 hours



	C _{max} (µg/mL)	AUC (µg.hr/mL)	T _{half} (hr)
rhIL-2	5.77	1.07	0.28
MDNA11	23.02	182.3	6.83

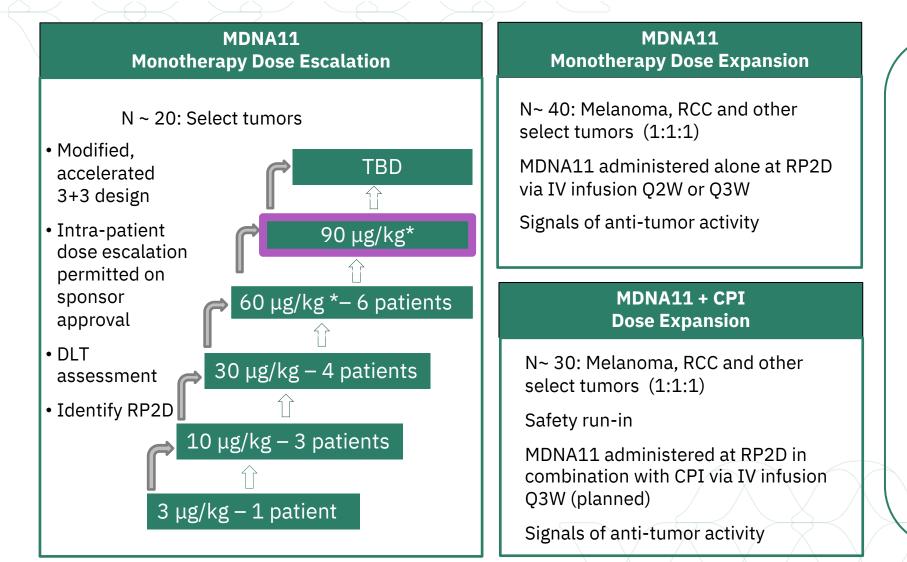
Naive C57Bl/6 mice IV dosed at 1 mg/kg IV

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MDNA11 Localization in CT26 Tumor Bearing Mice MDNA11 labelled w/VivoTag800; IV dose: 1 mg/kg; Tumor size: 150-200 mm³ ⁹

Phase 1/2 ABILITY Study Schema: Enrolling DL5



Endpoints:

- Safety/Tolerability
- ORR (RECIST 1.1)
- Clinical Benefit Rate (CBR) (CR+PR+SD)
- Survival EPs (TTE Analysis): PFS/OS
- Disease Control Rate (DCR)
- Duration of Response (DoR)
- Time to Relapse (TTR)

PK and PD Assessment:

- Immune Cell Profiling (Blood)
- Serum Cytokines
- Multiplex Immunofluorescence (Paired tumor biopsies)

Protocol Version 5

 NanoString Gene Expression (Paired tumor biopsies)

*Step-up dosing utilized: two priming doses of $30\mu g/kg$ given before the target dose

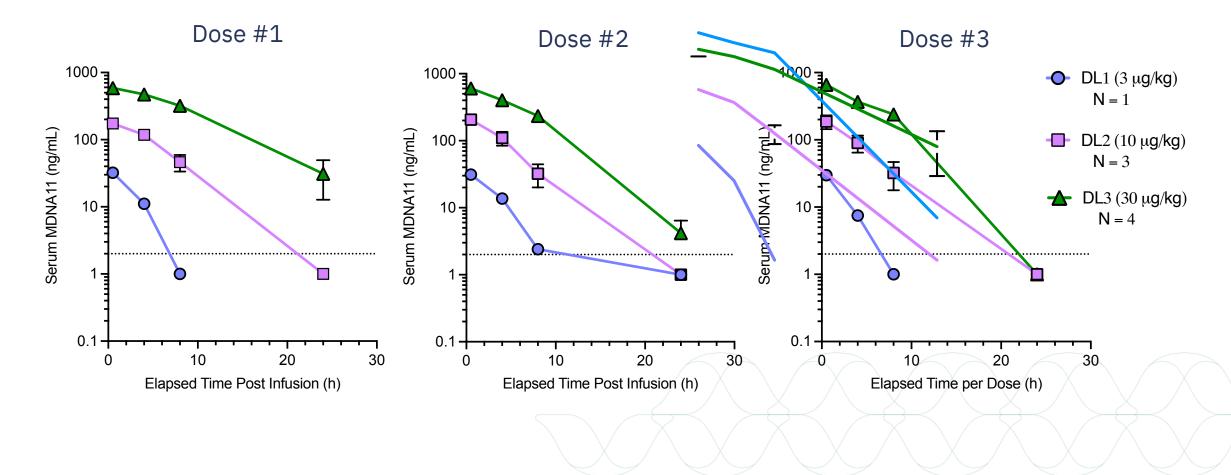
Patient Characteristics (N = 14, as of July 21, 2022)

Demographics/Performance	
Age (mean, yrs)	60.5
Male (%)	11/14 (79%)
Female (%)	3/14 (21%)
Baseline ECOG = 0	9/14 (64%)
Baseline ECOG = 1	5/14 (36%)
Primary Cancer Diagnosis	
Melanoma	7/14 (50%)
Renal Cell Carcinoma	1/14 (7%)
Pancreatic	2/14 (14%)
Sarcoma	2/14 (14%)
Squamous Cell Carcinoma	1/14 (7%)
Gastro-esophageal Adenocarcinoma	1/14 (7%)
Prior Systemic Therapies	
Lines of systemic therapies	1-4
Prior use of immunotherapy	11 /14 (79%)
Prior use of targeted therapy	4/14 (28%)
Prior use of chemotherapy	6/14 (43%)

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MDNA11 PK Profile in Patients with Advanced Solid Tumors

- > Highly similar PK plots following each of 3 repeat doses suggest no immunogenicity; ADA testing ongoing
- Dose-dependent increase in C_{max} and AUC

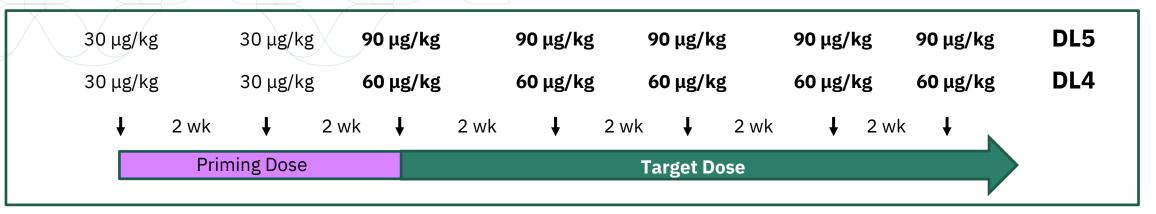


Summary of Key PK Parameters

	DL1: 3 μg/kg (N = 1)	DL2: 10 µg/kg (N = 3)	DL3: 30 μg/kg (N = 4)
Dose 1			
C _{max} (ng/mL)	32.2	174 ± 13.4	611 ± 188
AUC _{last} (h.ng/mL)	94.7	957 ± 187	5,497 ± 2,768
Dose 2			
C _{max} (ng/mL)	31.2	206 ± 40.5	604 ± 82.1
AUC _{last} (h.ng/mL)	130	969 ± 306	4,458 ± 1,373
Dose 3			
C _{max} (ng/mL)	29.8	188 ± 75	659 ± 259
AUC _{last} (h.ng/mL)	82.1	847 ± 387	3,434 ± 780

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Step Up Dosing (SUD) Strategy Implemented Starting at DL4



Objective:

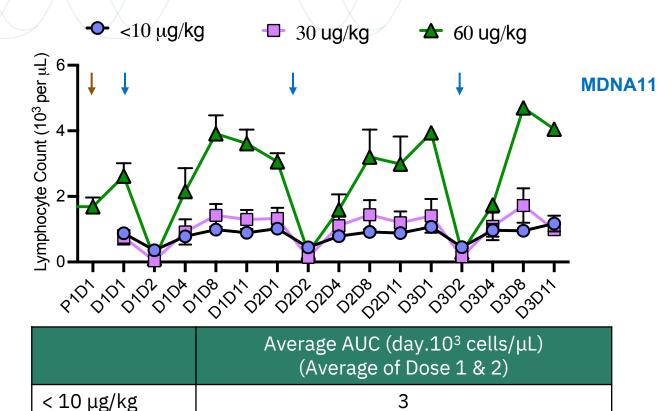
To pre-emptively mitigate the risk of potential toxicities during dose escalation in order to safely reach higher doses and achieve the best therapeutic response

Rationale:

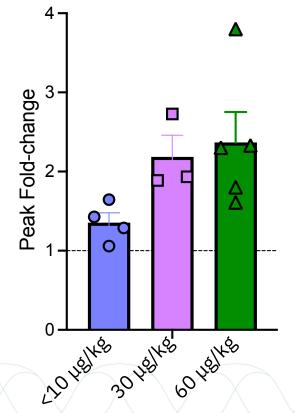
- SUD is adopted in clinical trials to deliver the highest achievable dose (Carlo-Stella et al., Hematological Oncology 2021; Trudel et al, Blood 2021; Hutchings et al., Blood 2020; Bartlett et al., ASCO 2019)
- A study in mice with MDNA11 demonstrated a > 2.5-fold increase in tolerability with SUD compared to a fixed dose administration

MDNA11 Induces Lymphocyte Expansion

Expansion of circulating lymphocytes irrespective of baseline count



Peak Fold-Change



Peak fold-change relative to baseline For < 10 μg/kg and 30 μg/kg, peak data for Dose 3 For 60 μg/kg, peak data for Target Dose 1

DL4 patients received 2 priming doses (30 µg/kg Q2W) prior to target dose (60 µg/kg Q2W))
AUC measured as area between minimum lymphocyte count values	

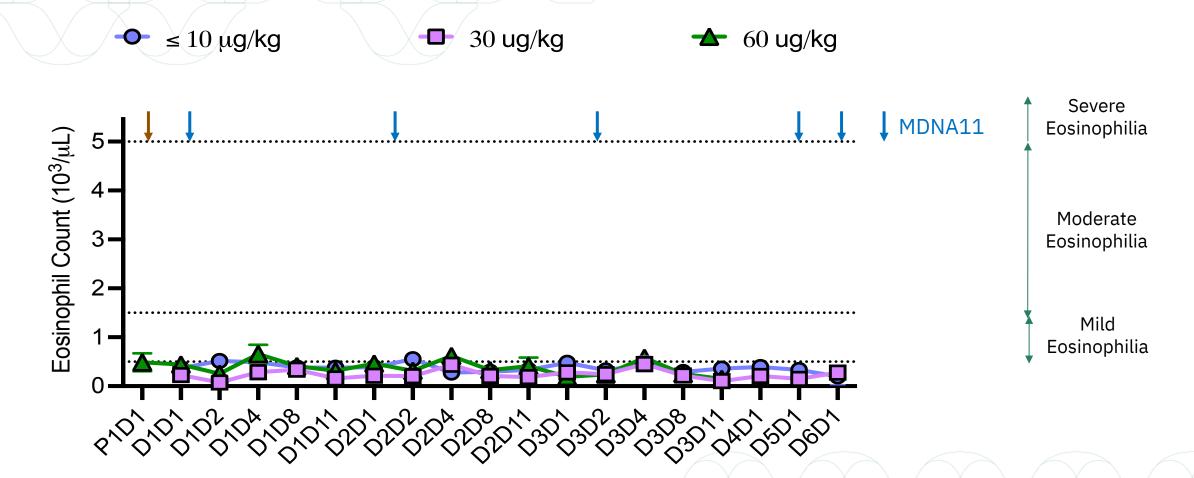
4.8

12.2

30 µg/kg

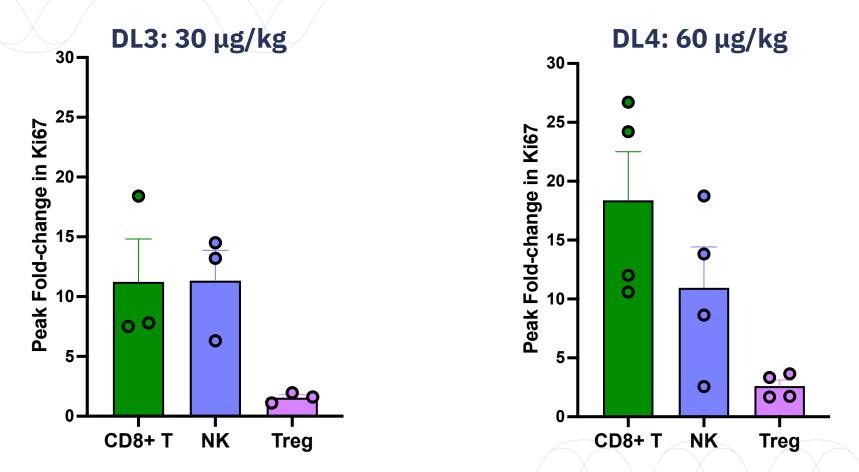
60 µg/kg

No Evidence of Eosinophilia (Associated with VLS)



DL4 patients received 2 priming doses (30 µg/kg Q2W) prior to start of target dose of 60 µg/kg (Q2W)

MDNA11 Preferentially Stimulates CD8⁺T & NK Cell Proliferation (Ki67)



Peak fold-change relative to respective baseline (D1D1 for DL3; P1D1 for DL4) DL4 patients received 2 priming doses (30 µg/kg Q2W) prior to start of target dose of 60 µg/kg (Q2W) DL3 data based on 3rd dose cycle

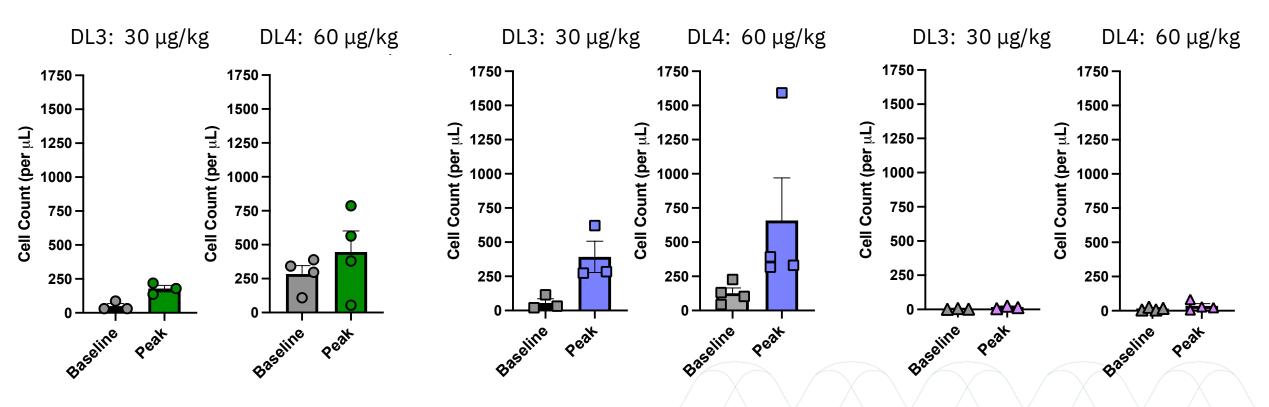
MDNA11 Preferentially Expands CD8⁺T and NK Cells Over Tregs

Increase in number of CD8+ T and NK cells with minimal change in Treg counts

CD8+ T Cells

NK Cells

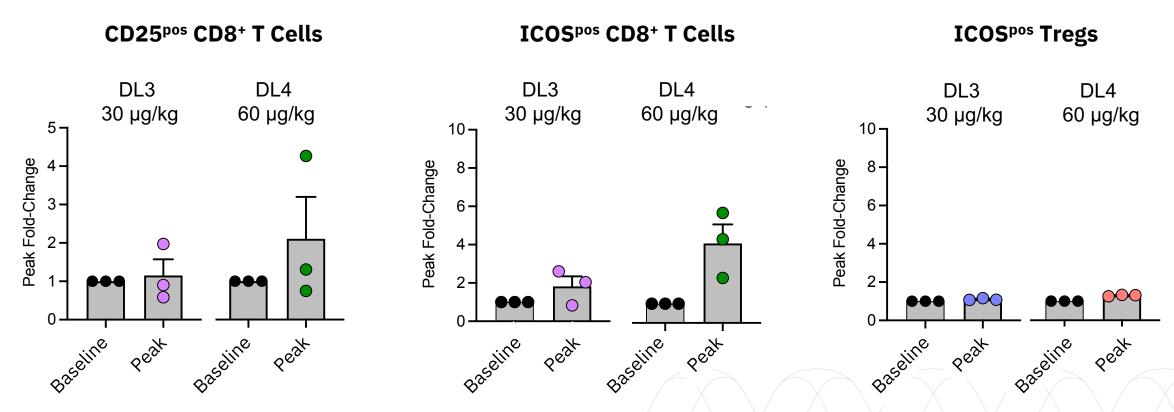
Tregs



DL4 patients received 2 priming doses (30 μ g/kg Q2W) prior to start of target dose of 60 μ g/kg (Q2W). DL3 data based on 3rd dose cycle; DL4 data based on 1st dose

CD8⁺ T Cell Activation Without ICOS Induction on Tregs

- Upregulation of CD25 and ICOS indicate CD8+ T cell activation
- > High dose rhIL-2 stimulates ICOS expression on Tregs and is associated with lack of therapeutic response*

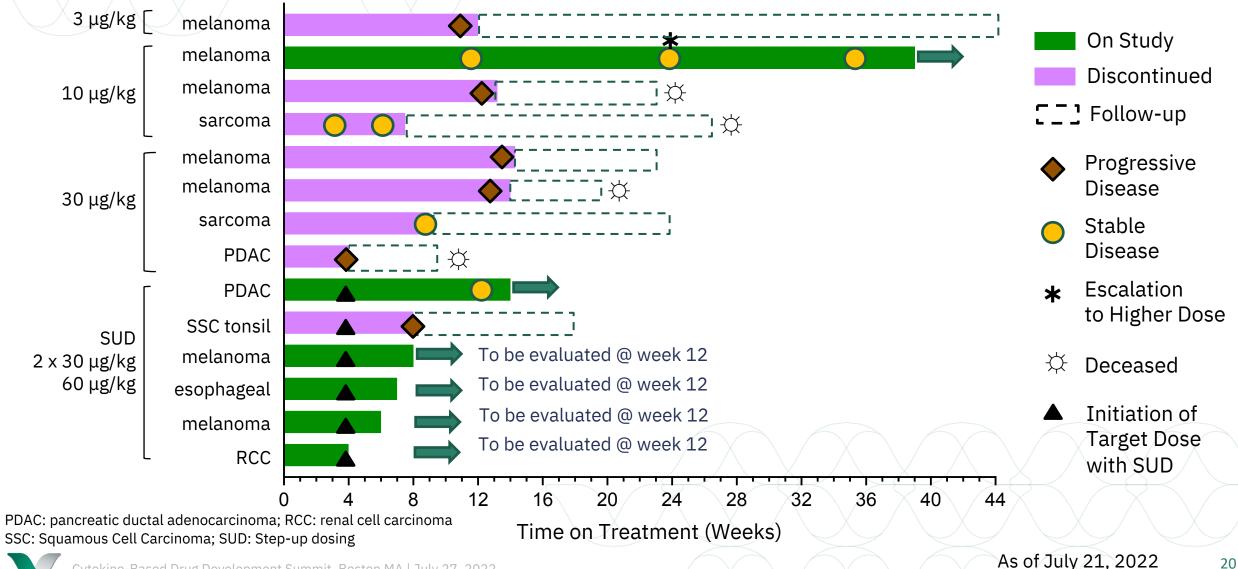


DL4 patients received 2 priming doses (30 µg/kg Q2W) prior to start of target dose of 60 µg/kg (Q2W). DL3 data based on 3rd dose cycle; DL4 data based on 1st dose Relative to baseline (D1D1 for DL3; P1D1 for DL4)

(*Sim et al., J Clinical Investigation, 2014)

Duration of Treatment and Summary of Response

Tumor Control Observed in 4 of 10 Evaluable Patients

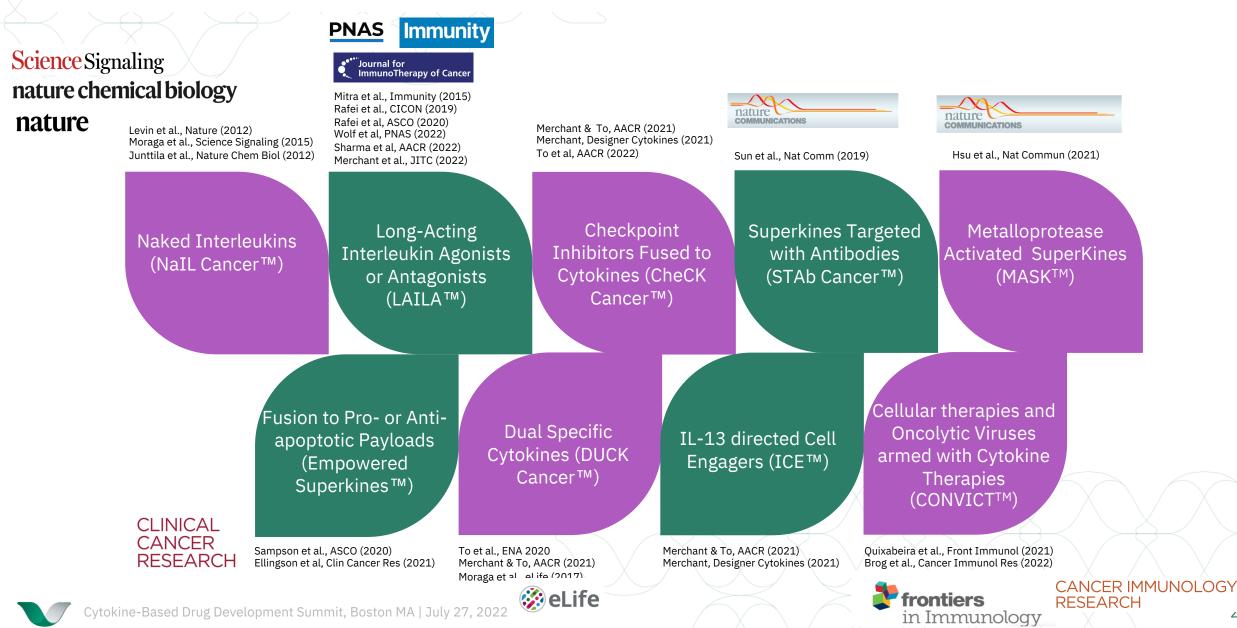


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MDNA11 is a Potentially Best-in-Class IL-2 Therapy

"Beta-only" IL-2 Super-Agonist	 High affinity for IL-2Rβ (CD122) and lack of affinity for IL-2Rα (CD25) enhances therapeutic efficacy and improves safety profile Competing IL-2 agents have comparatively low affinity for IL-2Rβ 	
Strong CD8+ T-cell and NK Cell Expansion with Clean Safety	 ~7.5 fold more potent on effector T cells and ~28 fold less potent on T_{regs} when compared to native IL-2 Preliminary Phase 1/2 data demonstrated preferential stimulation of CD8⁺ T and NK cells, with acceptable toxicity (no DLTs, no evidence of cytokine release syndrome or vascular leak syndrome) 	
Extended Half-Life	 Fusion to albumin scaffold extends half-life that allows for a highly convenient dosing schedule (Q2W) Albumin fusion provides additional benefit for localizing MDNA11 at the tumor site and tumor draining lymph nodes 	
Synergistic with Immune Checkpoint Inhibition	 Induced 100% tumor regression in preclinical models when co-administered with anti-CTLA4 and anti-PD1 Strong memory response with low immunogenicity risk, both as a monotherapy and in combination with anti-CTLA4 and anti-PD1 	

IL-2 Superkines Independently Validated Across Diverse Treatment Modalities



ACKNOWLEDGEMENTS

- > We are very grateful to the patients enrolled in the study and their friends and family
- > All Investigators and their teams for exceptional support at:
 - > Gallipoli Medical Research Foundation (GMRF), Brisbane, Australia
 - Chris O'Brien Lifehouse (COBLH), Sydney, Australia
 - > Scientia, Sydney, Australia
 - > ICON, Brisbane, Australia
 - Princess Margaret Hospital, Toronto, Canada

Thank You

