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Evolutionary IL-2 Superkines: *Past, Present and Future*

Next-Gen Cytokine Therapeutics Summit

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The PAST: Proleukin®



Validated with Durable Responses: Severe Toxicities Limit Use

High Unmet Need for Better IL-2 Immunotherapies



Short t 1/2



600,000 or 720,000 IU/kg IV Q 8 hours for up to 14 doses

Metastatic Renal Carcinoma



Metastatic Melanoma



- ORR = 14% (90% CI 10-19%)
- CR Rate = 5% (12/255)
- PR Rate = 9%
- Severe acute toxicities observed

Fyfe et al. (J Clin Oncol, 1995)

- ORR = 16% (90% CI 12-21%)
- CR Rate = 6% (17/270)
- PR Rate = 10% (26/270)
- Severe acute toxicities observed

Atkins et al. (J Clin Oncol, 1999)





The PRESENT: Superkines



Directed Evolution + Yeast Display = Tunable Superkines

Platform has generated extensive library of IL-2, IL-4, and IL-13 Superkines with unique properties



Basis of Versatile Superkine Platform

Merchant et al., ENA (2020) Levin et al., Nature (2012) Rafei et al, ASCO (2020) Moraga et al., Science Signaling (2015) Rafei et al., CICON (2019) Junttila et al., Nature Chem Biol (2012) Mitra et al., Immunity (2015) Naked Interleukins (NaIL Cancer[™]) (LAILA[™]) Cell therapies Armed Fusion to Pro- or Antiwith Superkines And apoptotic Payloads Viruses Armed with (Empowered superkines Superkines[™]) (CASAVA[™])

Sampson et al., ASCO (2020) Ellingson et al, Clin Cancer Res (2021)

Quixabeira et al., Front Immunol (2021)¹



Evolution of IL-2 Superkines: The MDNA109 Platform





MDNA109: A First-Generation IL-2 Super-agonist Without CD25 Dependency



MDNA109 Armed Virus Induces Potent Anti-Tumor Transcriptomic Program in TILs*



Quixabeira et al., Frontiers in Immunology 2021



Repression of Immune Suppression Genes



Work conducted independently of Medicenna at the University of Helsinki

Evolution of MDNA109 Superkine





Receptor Selectivity of MDNA109 Superkine Platform



	K _D [CD25 (IL-2Rα)]	K _D [CD122 (IL-2Rβ)]
IL-2 ^a	24 nM	210 nM
MDNA109 (1 st Gen.) ^a	26 nM	1.8 nM
MDNA109-Fc (2nd Gen.)b	14 nM	2.7 nM
MDNA109-Alb (<i>2nd Gen.)</i> a	56 nM	3.5 nM
MDNA19 (<i>3rd Gen.</i>) ^b	No binding	2.1 nM
MDNA11 (<i>3rd Gen.</i>) ^a	No binding	6.6 nM

a. BLI/Octet; b. SPR data

MDNA11 & MDNA19 Preferentially Stimulate Immune Effector Cells Over T_{regs}

STAT5 signaling in PBMCs from healthy donors





Protein	EC ₅₀ (nM)
rhIL2 (N = 4)	3.39
MDNA19 (N = 7)	0.37
MDNA11 (N = 3)	0.46

Protein	EC ₅₀ (nM)
rhIL2 (N = 4)	0.2
MDNA19 (N = 7)	0.071
MDNA11 (N = 3)	0.069

Protein	EC ₅₀ (nM)
rhIL2 (N = 4)	0.0056
MDNA19 (N = 7)	0.135
MDNA11 (N = 3)	0.160



Merchant et al, ENA 2020

MDNA109FA-Fc Exhibits Superior Anti-Tumor Efficacy in B16F10 Tumor Model







nature

Sun et al., Nat Comm., 2019

C57BL/6 mice (n = 5/group) were injected SQ with 5 \times 10⁵ B16F10 and treated IT with 5 µg of WT IL2-Fc or MDNA109FA-Fc on days 9, 12 & 15.

Work Conducted independently of Medicenna at the Chinese Academy of Sciences & University of Texas Southwestern Medical Center

MDNA11 Induces Preferential Tumor Infiltration of Effector Immune Cells



	CD8/Treg Ratio	
	Control	MDNA11
3 Days Post Dose	2.7	5.5
6 Days Post Dose	3.9	16.9

	NK/Treg Ratio	
	Control	MDNA11
3 Days Post Dose	3.8	15.9
6 Days Post Dose	4.8	8.4

B16F10 Tumor Model

Tumor bearing mice treated with a single dose of MDNA11 (5 mg/kg; IP)

MDNA11 + Anti-CTLA4 Induces Tumor Clearance, Protects Against Re-Challenges





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• All mice boosted with CT26 cells 5 days prior to analysis

MDNA11 Expands Lymphocytes Without Boosting Eosinophils in NHP



- Up to 9-fold increase in lymphocytes compared to pre-treatment.
- No expansion of eosinophils, potentially associated with VLS
- No evidence of Pulmonary Edema and no VLS observed
- No hypotension and no evidence of cytokine release syndrome nor ADA



MDNA11-01: ABILITY Phase 1/2 Study in Progress

Basket, accelerated sequential dose escalation and expansion study of MDNA11





The FUTURE <u>Bifunctional SuperKine</u> <u>ImmunoTherapies</u>

BiSKITs™



Potential of Medicenna's BiSKITs™ Platform



Work conducted independently of Medicenna at: ¹the Chinese Academy of Sciences; ²University of Texas Southwestern Medical Center

MDNA109 STABTM : Tumor Accumulation Enhances Response*





Left tumor: MC38 Right tumor: MC38-EGFR5

C57BL/6 mice were (n = 5/group) injected subcutaneously with 5 × 10⁵ of MC38-EGFR5 cells, and then i.v. treated on days 7 and 10 with PBS, 25 μ g of α EGFR-MDNA109FA or 25 μ g α TA99-MDNA109FA.



C57BL/6 mice (n = 5/group) were injected SQ with 5 × 10⁵ of B16F10-EGFR5 cells and IP treated with 25 µg of α EGFR-MDNA109FA-Fc or/and intratumorally treated with 50 µg of anti-PD-L1 on days 8, 11, and 14

Sun et al., Nat Comm., 2019

MDNA132 - Engineered Human IL-13 Targeting Tumor Specific Antigen (IL-13Rα2)



Moraga et al., Science Signaling, 2015

~4000 fold Selectivity for IL-13R α 2

SPR data K _D (nM)	IL-13Rα1	IL-13Rα2
IL-13	4.38	0.001
MDNA132	1600	0.0001

Tumors over-expressing IL-13Rα2 Bladder Cancer Colorectal Cancer **Pancreatic Cancer** Triple Negative Breast Cancer Glioblastoma Lung Cancer Head & Neck Cancer **Ovarian Cancer Prostate Cancer Mesothelioma**





Barderas et al., Cancer Res, 2012



MDNA19-413 is a Bi-functional Superkine

Targeting immunologic 'cold tumors' by modulation of TME



'Cold' tumors are not responsive to immunotherapies because of a Th2 (pro-tumoral) microenvironment:

- ➢ Low CD8⁺ & NK cell counts; high T_{reg} counts
- High number of immune-suppressive myeloid cells (TAM & MDSC)
- > TAMs & MDSCs overexpress the Type II IL-4 receptor

MDNA19-413 Inhibits IL-4 & IL-13 Induced Signaling & M2a Polarization



Concentration (nM)

Potential of Medicenna's Versatile Superkine Pipeline







Thank You!

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