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MT-6402 is a PD-L1 targeted engineered toxin body (ETB) composed of (Figure 1):

- De-immunized Shiga-like Toxin A subunit (SLTA) genetically fused to PD-L1 targeting antibody binding domain (scFv)
- HLA-A*02 restricted pp65 cytomegalovirus (CMV) antigen

FIGURE 1:
MT-6402

FIGURE 1:
MT-6402
structure

CMV pp65 peptide antigen

	Patient ID	Disease	Year of Birth	Sex	Prior CPI	HLA-A*02/CMV IgG positive	PD-L1 Assay	Result
Cohort 1 (16µg/kg)	1008-001	NSCLC	1945	M	Yes	Yes Yes	22C3	TPS 80%
	1004-002	NSCLC	1939	F	Yes	No	22C3	TPS 70%
	1001-001	Melanoma	1988	M	Yes	No	SP263	0.5% IC
	1002-003	Ovarian	1958	F	No	No	22C3	CPS > 1
	1005-002	Solid tumor	1974	M	No	No	22C3	TPS 10%
	1004-003	NSCLC	1958	M	Yes	Yes Yes	22C3	CPS > 1
Cohort 2 (24µg/kg)	1007-005	Esophageal	1951	M	Yes	Yes No	22C3	CPS 10
	1004-004	Solid tumor	1950	M	No	HLA TBD Yes	22C3	TPS 20%
	1001-002	NSCLC	1955	M	Yes	Yes No	22C3	TPS 10%
	1001-004	RCC	1971	F	Yes	Yes No	22C3	TPS 1%
	1008-002	Pancreatic	1960	M	No	No	SP142	5%
	1001-005	Skin SCC	1957	M	Yes	Yes Yes	22C3	CPS 3

	AE*	Grade	Comment
Cohort 1 (16µg/kg)	Anemia	3	Patient entered study with Grade 2 anemia
	Back pain	3	During infusion; treatment restarted within 30min after event resolved on Demerol and Phenergan; same patient had prior Grade 2 IRR
	Anorexia	2	
	CRS (SAE)	2	Recovered within 2 days
	Fever	2	
	IRR	2	Recovered within 1 hour
	Pruritus	2	
	Nausea	2	
Cohort 2 (24µg/kg)	Rash	3	Improved within 1 day on systemic steroids
	Fever	2	

Cohort 1

CD14+ Monocytes %change to baseline

Legend for Cohort 1:

- 1001-001 (non HLA-A*02, CMV+)
- 1002-003 (unknown)
- 1004-002 (non HLA-A*02, CMV+)
- 1004-003 (HLA-A*02, CMV+)
- 1008-001 (HLA-A*02, CMV+)

Cohort 2

CD14+ Monocytes %change to baseline

Legend for Cohort 2:

- 1001-002 (HLA-A*02, CMV-)
- 1001-004 (HLA-A*02, CMV-)
- 1001-005 (HLA-A*02, CMV+)
- 1004-004 (HLA unknown, CMV-)
- 1007-005 (HLA-A*02, CMV-)
- 1008-002 (non HLA-A*02, CMV-)

MDS (Freq of HLA+ cells)

Frequency of MDS

CD11b-HLA-D8 low/CD14+
 1001-001
 1001-003
 1005-002
 1008-001

Tregs (As a freq of CD4+ cells)

Frequency of Tregs

CD11b-HLA-D8 low/CD14+
 1001-001
 1001-003
 1005-002
 1008-001

Legend:

- 1001-001 (non HLA-A*02:01 CMV+)
- 1001-003 (unknown)
- 1005-002 (non HLA-A*02:01 CMV+)
- 1008-001 (HLA-A*02:01 CMV+)

MDS gating strategy: CD11b-HLA-D8 low/CD14+

Figure 3: CMV-specific CD8 T cell responses in 1008-003

Top Left: Percent of CMV-specific CD8 T cells (as a freq of CD8+ T cells)

CD8+ T cell	1008-003: Total CD8	1008-003: CMV-specific CD8
CD8+ T cell	~0.5	~0.5
CD8+ T cell + CMV	~2.5	~0.5
CD8+ T cell + CMV + IL-2	~1.5	~0.5
CD8+ T cell + CMV + IL-2 + IL-15	~0.5	~0.5
CD8+ T cell + CMV + IL-2 + IL-15 + IL-18	~0.5	~0.5
CD8+ T cell + CMV + IL-2 + IL-15 + IL-18 + IL-27	~0.5	~0.5
CD8+ T cell + CMV + IL-2 + IL-15 + IL-18 + IL-27 + IL-35	~2.5	~0.5

Top Right: Total CD8 and CMV-specific CD8 Fold Change

CD8+ T cell	1008-003: Total CD8	1008-003: CMV-specific CD8
CD8+ T cell	1.0	1.0
CD8+ T cell + CMV	~1.5	~1.0
CD8+ T cell + CMV + IL-2	~1.0	~1.0
CD8+ T cell + CMV + IL-2 + IL-15	~1.0	~1.0
CD8+ T cell + CMV + IL-2 + IL-15 + IL-18	~1.0	~1.0
CD8+ T cell + CMV + IL-2 + IL-15 + IL-18 + IL-27	~1.0	~1.0
CD8+ T cell + CMV + IL-2 + IL-15 + IL-18 + IL-27 + IL-35	~1.5	~1.0

Bottom Left: Percent of CMV-specific CD8 T cells (as a freq of CD8+ T cells)

CD8+ T cell	1008-003: Total CD8	1008-003: CMV-specific CD8
CD8+ T cell	~1.0	~1.0
CD8+ T cell + CMV	~1.8	~1.0
CD8+ T cell + CMV + IL-2	~1.0	~1.0
CD8+ T cell + CMV + IL-2 + IL-15	~1.0	~1.0
CD8+ T cell + CMV + IL-2 + IL-15 + IL-18	~1.0	~1.0
CD8+ T cell + CMV + IL-2 + IL-15 + IL-18 + IL-27	~1.0	~1.0
CD8+ T cell + CMV + IL-2 + IL-15 + IL-18 + IL-27 + IL-35	~1.0	~1.0

Bottom Right: Total CD8 and CMV-specific CD8 Fold Change

CD8+ T cell	1008-003: Total CD8	1008-003: CMV-specific CD8
CD8+ T cell	1.0	1.0
CD8+ T cell + CMV	~1.5	~1.0
CD8+ T cell + CMV + IL-2	~1.0	~1.0
CD8+ T cell + CMV + IL-2 + IL-15	~1.0	~1.0
CD8+ T cell + CMV + IL-2 + IL-15 + IL-18	~1.0	~1.0
CD8+ T cell + CMV + IL-2 + IL-15 + IL-18 + IL-27	~1.0	~1.0
CD8+ T cell + CMV + IL-2 + IL-15 + IL-18 + IL-27 + IL-35	~1.0	~1.0

	Cytokines*	PBMC Changes	
All Patients (Irrespective of HLA or CMV status)	▲ CCL2/MCP-1	▼ Monocytes ▼ MDSCs	Movement of myeloid cells to tissues and/or direct cell kill
	▲ IL-2	▲ Tbet+ CD8 T cells ▲ NK cells	
AST-Engaged Only (HLA-A*02, CMV+)	▲ TNFα ▲ IL-10	▲ Peripheral CD8 T cells ▲ Peripheral CMV-specific T cells (initial expansion)	Checkpoint break plus AST driven activation of CMV-specific T cells
	▲ IP-10/CXCL10	▼ Peripheral CMV-specific T cells (after first dose) ▼ Peripheral dendritic cells	Migration of dendritic cells and CMV-specific T cells from periphery

Figure 2 consists of three line graphs showing the effect of treatment cycles on HIV-1 RNA levels, Mo-MSDC frequency, and CMV-specific CD8+ T-cell frequency.

Left Graph: Gradual increase in monocyte recovery between doses starting at Cycle 6

This graph shows the percentage change in monocyte counts from baseline over 24 cycles. The y-axis ranges from -100% to 50%. The x-axis shows cycles from C1D1-Pre to C24D1-Pre. A dashed blue line indicates a gradual increase in monocyte recovery starting at Cycle 6, reaching approximately 10% by Cycle 24.

Middle Graph: Rebound in MSDSCs starting at Cycle 5

This graph shows the frequency of Mo-MSDC over 24 cycles. The y-axis ranges from 0 to 6. The x-axis shows cycles from C1D1-Pre to C24D1-Pre. A dashed blue line indicates a rebound in Mo-MSDC frequency starting at Cycle 5, reaching approximately 5.5 by Cycle 24. A blue star is marked at Cycle 5, and a blue circle highlights the rebound period from Cycle 5 to Cycle 24.

Right Graph: Rebound in peripheral CMV-specific CD8+ T-cells starting at Cycle 5

This graph shows the frequency of CMV-specific CD8+ T-cells over 24 cycles. The y-axis ranges from 0.0 to 2.5. The x-axis shows cycles from C1D1-Pre to C24D1-Pre. A dashed blue line indicates a rebound in CMV-specific CD8+ T-cell frequency starting at Cycle 5, reaching approximately 2.0 by Cycle 24. A blue star is marked at Cycle 5, and a blue circle highlights the rebound period from Cycle 5 to Cycle 24.

Note: these images are untouched and are presented exactly as provided to MTEM.

[illegible]