CD86: A Novel Prognostic Marker in Acute Lymphoblastic Leukemia Patients

Thesis

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μL : Micro liter

ALL : Acute lymphoblastic leukemia

AML : Acute myeloid leukemia
 APCs : Antigen presenting cells
 ARF : ADP-ribosylation factor

AUC : Area under the curve

B-ALL: B-cell Acute lymphoblastic leukemia

BCP-ALL: B-cell precursor ALL

BCR/ABL: Breakpoint cluster region/Abelson

BFM: Berlin-Frankfurt-Münster

BM : Bone marrow

CALLA: Common ALL antigen
CBC: Complete blood count
CCG: Children's Cancer Group
CD: Cluster of differentiation
CHL: Classic Hodgkin lymphoma

CLL: Chronic lymphocytic leukemia

C-Myc : Cellular myelocytoma gene

CNS : Central nervous system

CR : Complete remission

CSCs : Cancer stem cellsCSF : Cerebrospinal fluid

CTL : Cytotoxic T- lymphocytes

CTLA-4 : Cytotoxic T-lymphocyte antigen 4

DCs : Dendritic cells

DNA : Deoxyribonucleic acidEBF1 : Early B cell factor 1

EDTA : Ethylene-diamine-tetra-acetic acid

EMI : Extramedullary involvement

FAB: French-american-british

FCM: Flow cytometry

FISH: Fluorescence in situ hybridization

FITC: Fluorescein isothiocyanate

FRET : Forster resonance energy transferGPI : Glycosyl phosphatidyl inositol

HA : Hyaluronic acidHb : Hemoglobin

HLA: Human leucocyte antigen

HLA-DR: Human leukocyte differentiation antigen

HS: Highly significant

HSCs: stem cells

iAMP21 : Intra-chromosomal amplification of the

chromosome 21

ICAM-1: Intercellular adhesion molecule-1

ICOS-L : Inducible costimulator ligand

IgC : Immunoglobulin constantIgE : Immunoglobulin type EIgG : Immunoglobulin type G

IGSF: Immunoglobulin superfamily

IgV : Immunoglobulin variable

IKZF1 : IKAROS family zinc finger

IL : Interleukin

IPMs: Immunophenotypic markers

IPT: Immunophenotyping

IS : Immunological synapse

kDa : Kilodalton

LSCs : Leukemic stem cells mCD86 : Membranous CD86

MFI: Mean florescence intensity

MHC: Major histocompatibility complex

MLL : Myeloid-lymphoid lineage

MM: Multiple myeloma

MoAbs : Monoclonal antibodies

MPO: Myelo-peroxidase

MRD: Minimal residual disease

MS : Multiple sclerosis

NCI : National Cancer Institute

NK: Natural Killer cell

NPV : Negative predictive value

NS : Non significant

p : Short arm of chromosome

P : Probability
PAX5 : Paired box 5

PB: Peripheral blood

PBS : Phosphate buffered salinePCR : Polymerase chain reaction

PD-L1 : Programmed cell death-1 ligandPD-L2 : Programmed cell death-2 ligand

PE: Phycoerythrin

PHF6: PHD finger protein 6

PLT : Platlets

PMNs : Polymorphonuclear cells
 POG : Pediatric Oncology Group
 PPV : Positive predictive value
 q : Long arm of chromosome

RNA : Ribonucleic acid

ROC : Receiver operating characteristic

RT-PCR: Reverse transcriptase polymerase chain reaction

RUNX1 : Runt related transcription factor 1

S : Significant

sCD44 : Serum levels of CD44
sCD86 : Serum levels of CD86
SD : Standard deviation

Sig : Significance

SLE : Systemic lupus erythematosis

t : Translocation

T-ALL: T-cell acute lymphoblastic leukemia

TCR : T-cell receptorTCR : T cell receptor

TdT: Terminal deoxynucleotidyl transferase

Teff: Effector T cells

TEL-AML1: Tel-acute myeloid leukemia

Th1 : T helper type 1Th2 : T helper type 2

TIL: Tumor infiltrating lymphocytes

TLC: Total leukocytic count

TNFR: Tumor necrosis factor receptor

TNFRSF: Tumor necrosis factor superfamily

Treg: Regulatory T cell

VISTA: V-domain immunoglobulin suppressor of T cell

activation

WBC: White blood cell

WHO: World health organization

WT1 : Wilms tumor 1
X² : Chi- square test

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Introduction

Leukemia is a hematopoietic malignancy that results from the clonal proliferation of bone marrow cells with impaired differentiation, regulation, and cell death. The acute leukemias arise from neoplastic transformation of hematopoietic stem cells or progenitors with aberrant differentiation and proliferation (Gowda and Dovat, 2013).

Acute lymphocytic leukemia (ALL) is the most common hematological malignancy in childhood and accounts for about 20% of acute leukemia in adults (**Liew** et al., 2012).

On the basis of ontogenic classification, ALL is divided into T-lineage ALL and B-lineage ALL. B-lineage ALL is characterized by the expression of the B-cell markers CD19, CD22 and CD79a (**Pui et al., 2008**).

T-cell acute lymphoblastic leukemia (T-ALL) is a rare, aggressive malignancy of thymocytes and corresponds to a heterogeneous group of leukemia arrested at various stages of lymphoid development. T-ALL constitutes 15% of all childhood ALL and 25% of adult ALL; approximately 30% of patients relapse within the first year of treatment and the outcome is usually death (**Pui et al.**, **2004-a**).

Leukemogenesis is a multistep process that requires the accumulation of alterations in a hematopoietic progenitor cell at multiple stages. The leukemic stem cell (LSC) hypothesis postulates that leukemia are hierarchically organized and leukemic stem cells have the capacity to self-renew, give rise to more differentiated progeny and maintain the leukemia long-term (Magee et al., 2012).

CD86 is a member of B7 family, which consists of cell-surface proteins that regulate costimulatory or coinhibitory signals by binding to their ligands (**Greaves and Gribben, 2013**).

Recognition of CD86 ligand by co-stimulatory CD28 and co-inhibitory CTLA-4 receptors plays an important role in influencing immune responses by proliferation and suppression of effector T cells respectively (**Postow et al., 2015**).

CD86 is expressed on the surface of antigen presenting cells (APCs) as monocytes and dendritic cells (DCs) (Lim et al., 2012). Its expression was found to be associated with many hematological malignancies such as acute myeloid leukemia (AML) and it was reported as a marker of poor prognosis in it (Barrett and Le Blanc,

2010). The expression of it in ALL cells is variable but it is mostly associated with poor prognosis in those patients (Mansour et al., 2014).

Some of the most recent therapeutic developments for acute leukemia depend on the involvement of costimulatory pathways and molecules including CD86 (Couzin-Frankel, 2013).