

# بسم الله الرحمن الرحيم

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بقسم التوثيق الإلكتروني بمركز الشبكات وتكنولوجيا المعلومات دون أدنى مسئولية عن محتوى هذه الرسالة.

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AIN SHAMS UNIVERSITY

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# CLINICAL OUTCOMES FOR PATIENTS WITH ACUTE MYELOID LEUKEMIA HARBORING IDH'I MUTATION AFTER INTENSIVE CHEMOTHERAPY

#### Thesis

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### **List of Abbreviations**

Al	bb.		Full term
•	2HGD	•	2-hydroxyglutarate dehydrogenases
•	AML	•	Acute myeloid leukemia
•	APL	•	Acute promyelocytic leukemia
•	ASXL1	•	Additional sex comb-like 1 mutations
٠	ATO	•	Arsenic trioxide
•	ATRA	•	All-trans retinoic acid
•	CBFA2	•	Core-binding factor subunit-2
•	CBFB	•	Core binding factor beta subunit
•	CD	•	Common differentiation
•	CEBPA	•	Ccaat enhancer binding protein
•	CIBMTR .	•	Center of international blood and marrow
			transplant research
•	CN-AML.	•	Cytogenetically normal
•	CR	•	Complete remission
•	DFS	•	Disease free survival
•	DIC	•	Disseminated intravascular coagulation
•	DNMT3A	•	Dna methyltansferase 3a mutations
•	ELN	•	European leukemianet
•	EVI1	•	Ecotropic virus integration site 1
•	FAB	•	French-american- british classification
•	FLT3	•	Fms-like tyrosine kinase 3 mutations
•	HLADR	•	Human leukocyte antigen-antigen d related
•	HMAs	•	Hypomethylating agents
•	HRM	•	High resolution melt

- HSCT ..... Hematopoietic stem cell transplantation IDHs..... Isocitrate dehydrogenases ITD..... Internal tandem duplications JAK2 ...... The janus kinase 2 gene JM ..... Juxta-membrane ■ L-2HG..... L-2-hydroxyglutarate enantiomer MDS ..... Myelodysplastic syndrome MLL ..... Mixed lineage leukemia MPD..... Myeloid proliferative disorder MPL ..... Myeloproliferative leukemia MRD ..... Minimal residual leukemia MYH11... Smooth muscle myosin heavy chain 11 NADPH... Nicotinamide adenine dinucleotide phosphate NGS..... Next generation sequencing NPM1..... Nucleophosmin 1 OS ..... Overall survival PCR ..... Polymerase chain reaction PHDs..... Prolyl hydroxylases PML-Promyelocytic leukemia/retinoic acid receptor RARA ..... alpha PTD ..... Partial in tandem duplications RUNX1... Runt-related transcription factor
  - Therapy-related amlTerminal deoxynucleotidyl transferase

RUNX1T1

SRSF2 .....

t-AML .....

TdT.....

Runx1 partner transcriptional co-repressor 1

Serine/ arginine-rich- splicing-factor-2 gene

- TET2 ...... Ten–eleven translocation 2
- TKD ....... Tyrosine kinase domain
- TKI..... Tyrosine kinase inhibitors
- UK NCRC United kingdom national cancer research institute
- WT1 ...... Wilms' tumor 1 gene
- α-KG ...... A-ketoglutarate

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#### INTRODUCTION

Acute myeloid leukemia (AML) is heterogeneous myeloid disorder with multifactorial pathogenic mechanisms and a broad range of prognosis. AML is characterized by clonal proliferation of poorly differentiated cells of the myeloid lineage (*Dohner et al.*, 2015).

The pathogenesis involve recurrent genomic alterations, including somatic gene mutations and/or chromosomal abnormalities, that can define biologically distinct clinical subtypes (*Vardiman et al.*, 2009).

Comprehensive genomic profiling at the time of diagnosis can inform disease classification, risk stratification and prognosis and ultimately allow for more selective therapeutic interventions. Alterations to cellular metabolism, as well as somatic mutations of genes essential to epigenetic regulation, are implicated in the pathogenesis of several human malignancies(*Cairns et al., 2011; Conway et al., 2014*).

Isocitrate dehydrogenases (IDHs) are homodimeric enzymes involved in diverse cellular processes, including adaptation to hypoxia, histone demethylation and DNA modification(*Clark et al.*, *2016*).

IDH1 protein catalyze the oxidative decarboxylation of isocitrate to  $\alpha$ -ketoglutarate ( $\alpha$ -KG) to produce reduced nicotinamide adenine dinucleotide phosphate.

Diverse dioxygenases depend on sufficient levels of  $\alpha$ -KG for multiple cellular processes, as well as for epigenetic regulation (*Molenaar et al.*, 2014).

IDH1 enzymes are localized to the cytoplasm and peroxisomes (*Clark et al.*, 2016).

Somatic mutations in IDH1 (mIDH1) genes have been described in both solid and hematological malignancies (*Stein*, 2016).

IDH1mutations are heterozygous, retaining one wild-type, suggestive of an oncogenic gain of function. IDH proteins are encoded by the IDH1 gene located at chromosome 2q33. Recurrent IDH1 mutations are missense variants leading to a single amino-acid substitution of arginine residues at codon 132 in exon 4 of the IDH1 gene. Additionally, a germline-synonymous single-nucleotide polymorphism (rs11554137) located in codon 105 in exon 4 of the IDH1 gene has been reported to have prognostic relevance in AML(*Willander et al., 2014; Wagner et al., 2010*). In our study we will provide an overview of the Clinical Outcomes for Patients with Acute Myeloid Leukemia Harboring IDH1 mutation after Intensive Chemotherapy.

#### **AIM OF THE WORK**

The aim of the present study is to detect IDH1 Mutation in adult Egyptian AML patient and find correlation between the mutation and prognosis & survival in those patients after intensive chemotherapy.

# Chapter (1) Acute Myeloid Leukemia

#### I. Introduction

Acute myeloid leukemia (AML) is a heterogeneous disorder characterized by clonal expansion of myeloid progenitors (blasts) in the bone marrow and peripheral blood. Previously incurable, AML is now cured in approximately 35%-40% of patients younger than age 60 years old. For those >60 years old, the prognosis is improving but remains grim. Studies have revealed that the disorder arises from a series of recurrent hematopoietic stem cell genetic alterations accumulated with age. Using deep sequencing techniques on primary and relapsed tumors, a phenomenon called clonal evolution has been characterized with both founding clones and novel subclones, impacting the therapeutic approach. Despite an increased understanding of AML biology, our efforts to this point in changing treatment strategy have been disappointing. In this review, we discuss the current diagnostic and prognostic strategies, current treatment approaches and novel therapies critical to AML management (Khwaja et al., 2016).

#### II. Morphology

Morphologically, AML blasts vary in size from slightly larger than lymphocytes to the size of monocytes or larger. The nuclei are large in size, varied in shape and usually contain several nucleoli. AML blasts express