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La terapia con anti-BCL2: nuovi target

INTRODUZIONE

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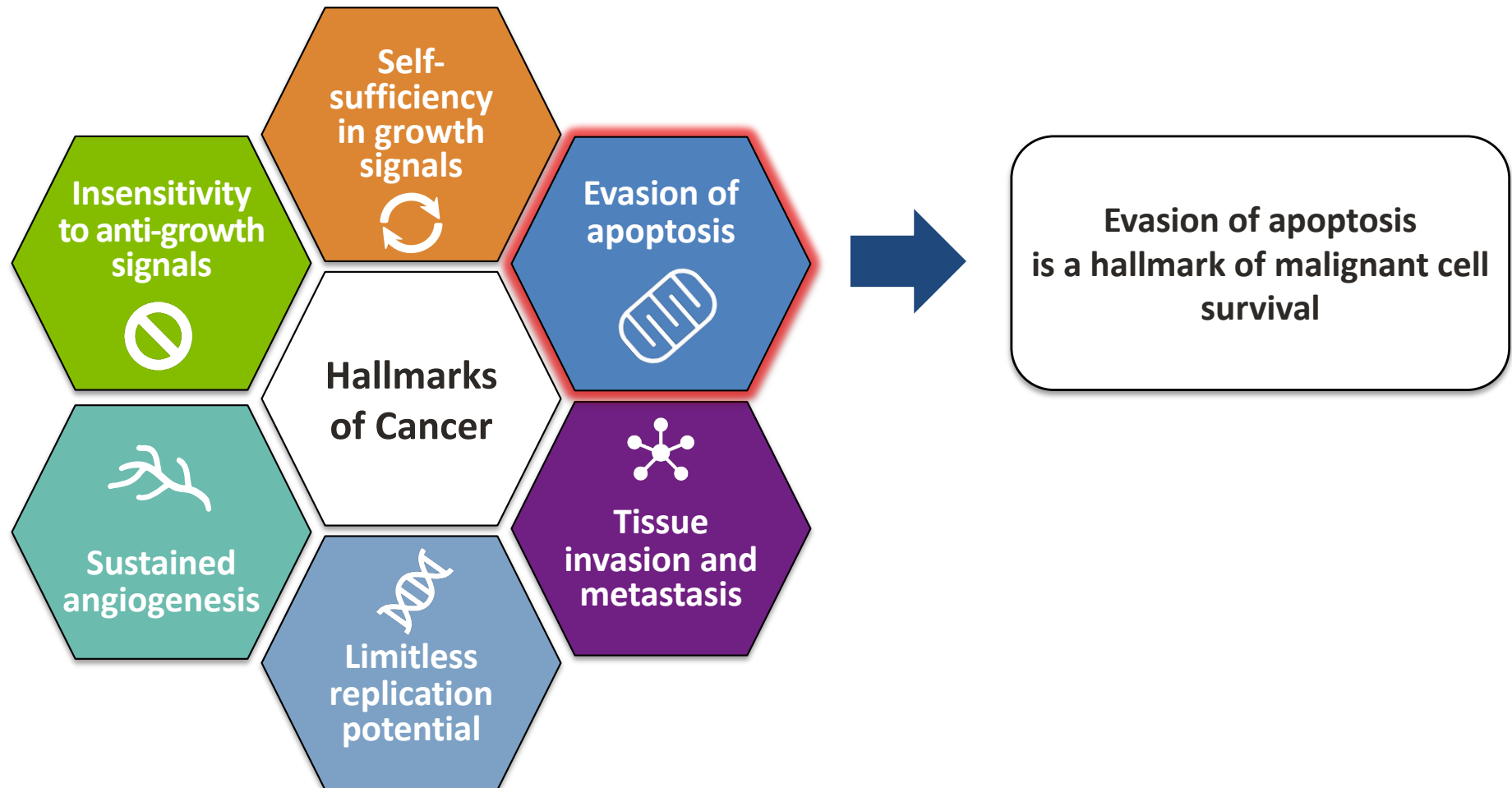
Marco Montillo

Conflict of Interest

- Abbvie
- Acerta/Astra Zeneca
- Gilead
- Janssen
- Roche
- Verastem

Cancer Pathogenesis Is Dependent on Numerous Mechanisms

Oncogenesis is driven by a complex interplay of genetic alterations and mechanisms that ensure cell survival and proliferation

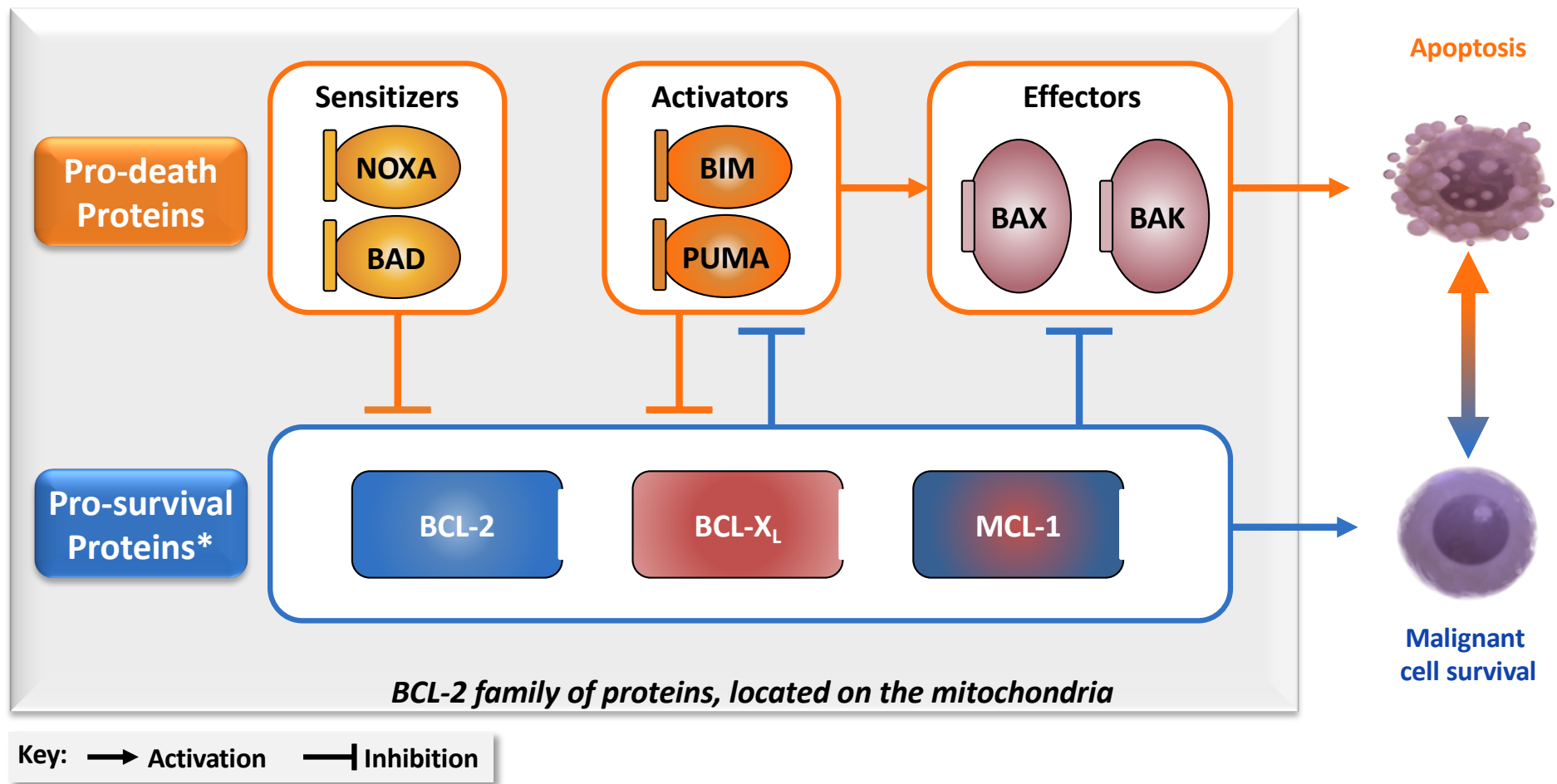


BCL2 family proteins

- Bcl2 was the first discovered regulator of apoptosis when it was found as translocated in patients suffering from follicular B cell lymphoma, 30 years ago.
- Nineteen proteins of the human 'Bcl-2 family' have been described.
- Seven members of this family (Bcl-2, Bcl-XL, Mcl-1, Bcl-B, Bcl-w and A1/Bfl1) are anti-apoptotic proteins while the rest (Bax, Bak, Bok, Bim, Bmf, Puma, Noxa, Bad, Bid, Bcl-XS, BiNP3 and Hrk) have a pro-apoptotic function.
- The balance between protein anti-apoptotic and pro-apoptotic proteins defines the fate of cell

Apoptosis Is Regulated by the BCL-2 Family of Proteins

BCL-2 family proteins include both pro-survival (anti-apoptotic) and pro-death (pro-apoptotic) proteins with opposing functions¹⁻⁴

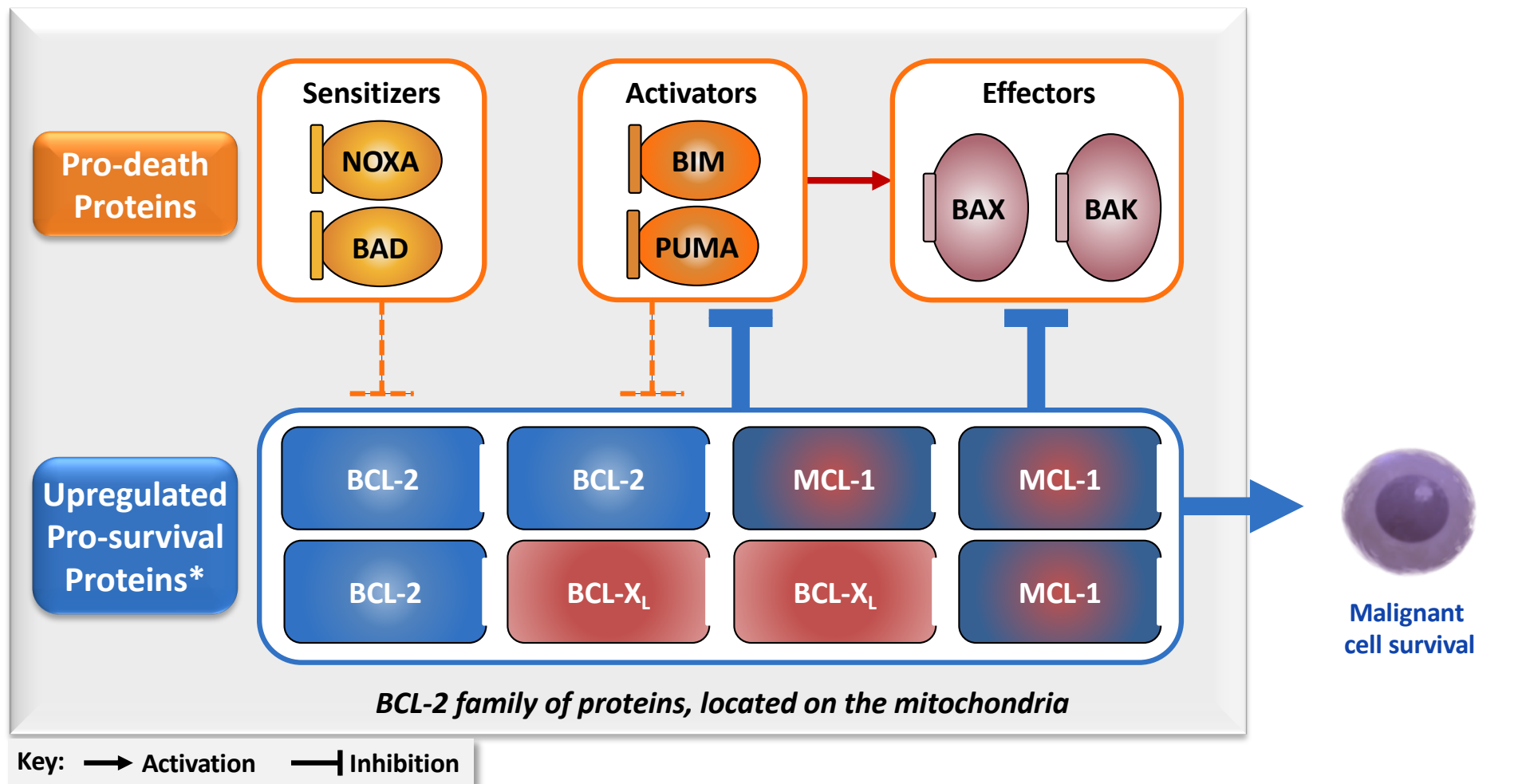


* Also includes: BCL-w and BFL-1/BCL2-A1.
BCL-2, B-cell lymphoma 2.

1. Plati J, et al. *Integr Biol (Camb)* 2011; **3**:279–296;
2. Czabotar PE, et al. *Nat Rev Mol Cell Biol* 2014; **15**:49–63;
3. Levenson JD, et al. *Cancer Discov* 2017; **7**:1376–1393;
4. Valentin R, et al. *Blood* 2018; **132**:1248–1264.

Malignant Cells Can Evade Apoptosis by Upregulating BCL-2 and Other Pro-survival Proteins

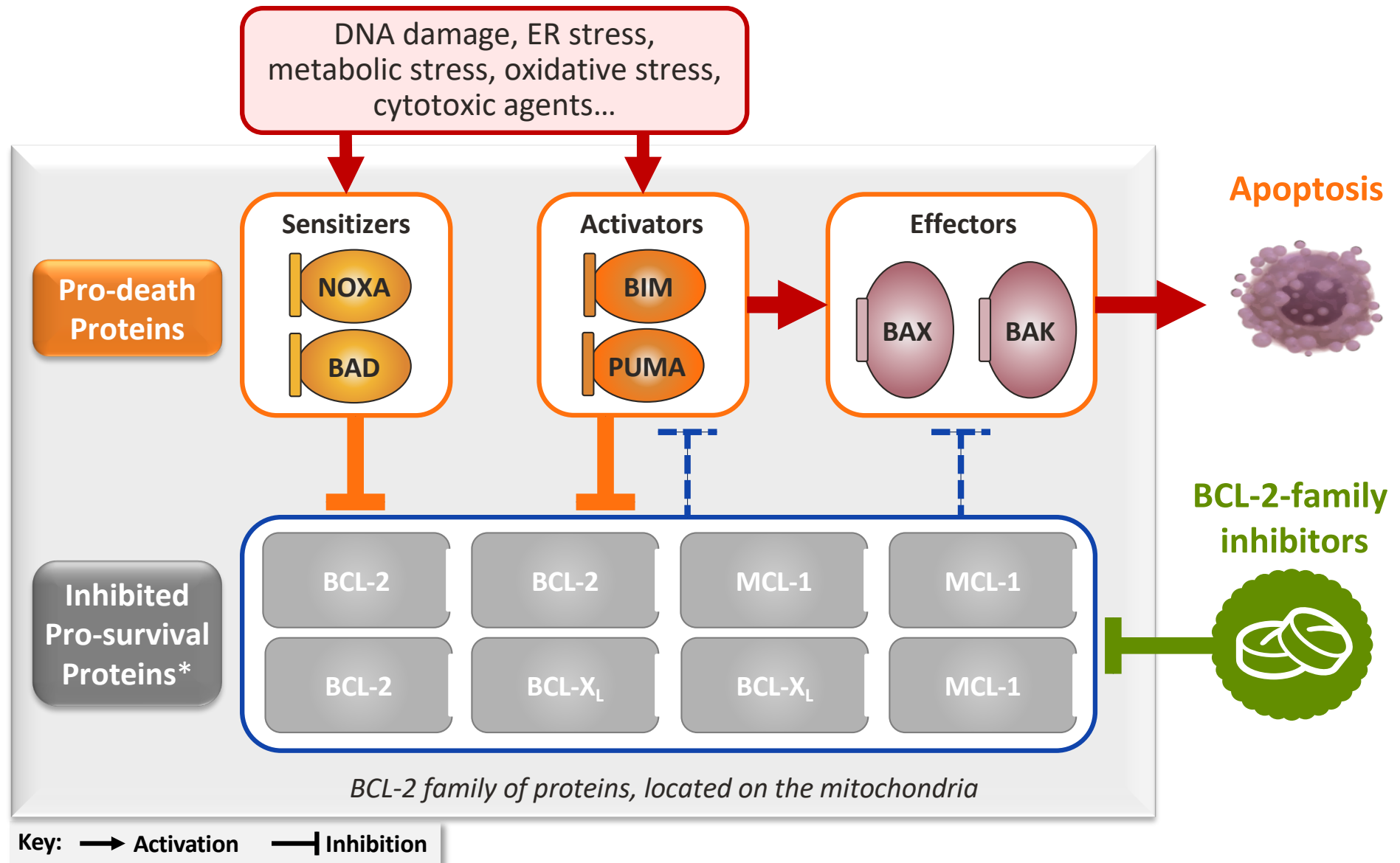
Malignant cells often evade apoptosis by upregulation of pro-survival proteins, such as BCL-2, MCL-1, and BCL-X_L



* Also includes: BCL-w and BFL-1/BCL2-A1.

1. Levenson JD, et al. *Cancer Discov* 2017; **7**:1376–1393;
 2. Czabotar PE, et al. *Nat Rev Mol Cell Biol* 2014; **15**:49–63; 3. Adams JM & Cory S. *Oncogene* 2007; **26**:1324–1337;
 4. Letai A, et al. *Cancer Cell* 2002; **2**:183–192; 5. Certo M, et al. *Cancer Cell* 2006; **9**:351–365.

Agents that Inhibit Pro-survival Proteins Can Promote Apoptosis in Malignant Cells



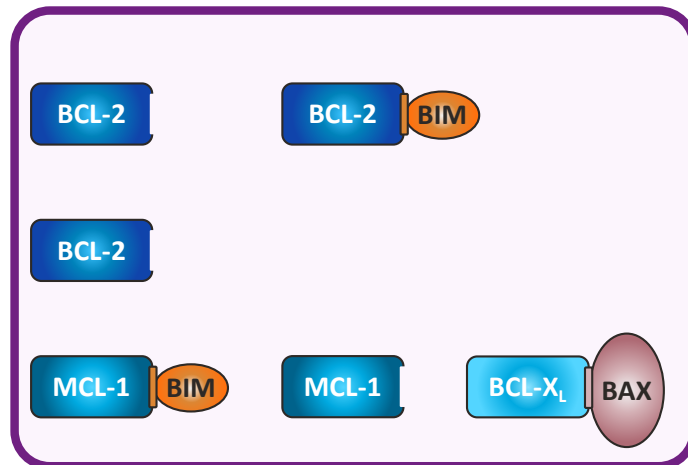
* Also includes: BCL-w and BFL-1/BCL2-A1.

1. Levenson JD, et al. *Cancer Discov* 2017; **7**:1376–1393;
 2. Czabotar PE, et al. *Nat Rev Mol Cell Biol* 2014; **15**:49–63; 3. Adams JM & Cory S. *Oncogene* 2007; **26**:1324–1337;
 4. Letai A, et al. *Cancer Cell* 2002; **2**:183–192; 5. Certo M, et al. *Cancer Cell* 2006; **9**:351–365.

Malignant Cells with High Dependence on the BCL-2 Protein for Survival Are “Primed for Death”

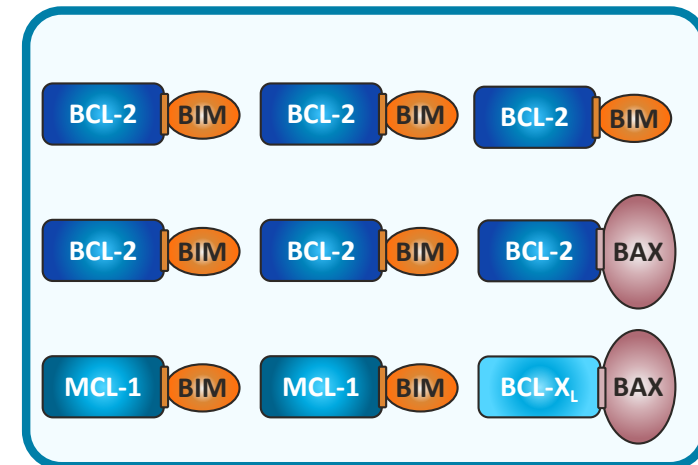
Unprimed healthy cell: High apoptotic threshold

- Low level of sequestered pro-death proteins
- Released pro-death proteins may be re-captured by available pro-survival proteins and may not be sufficient to trigger apoptosis





Primed malignant cell: Low apoptotic threshold

- High levels of pro-survival proteins and high levels of sequestered pro-death proteins
- Sufficient release of pro-death proteins is likely to initiate apoptosis
- “BCL-2 primed”: primed due to high levels of BCL-2 relative to other pro-survival proteins

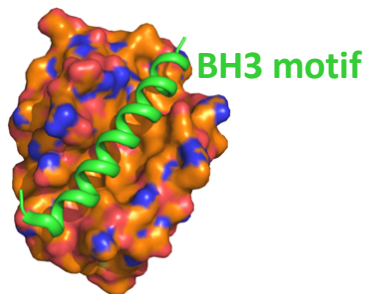
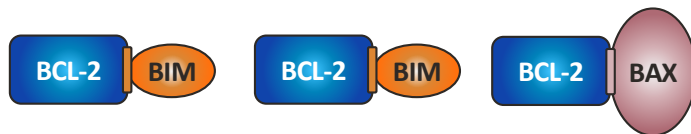


Key:

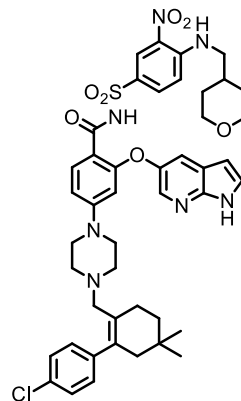
  Bound BIM–BCL-2

Venetoclax Is a Highly Selective, Potent, Oral BCL-2 Inhibitor Designed to Induce Apoptosis in Malignant Cells

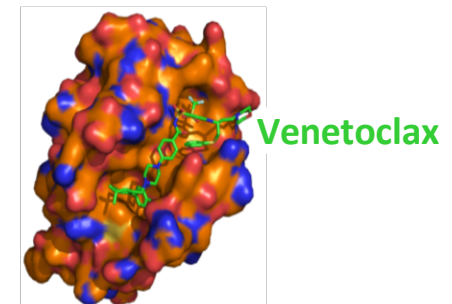
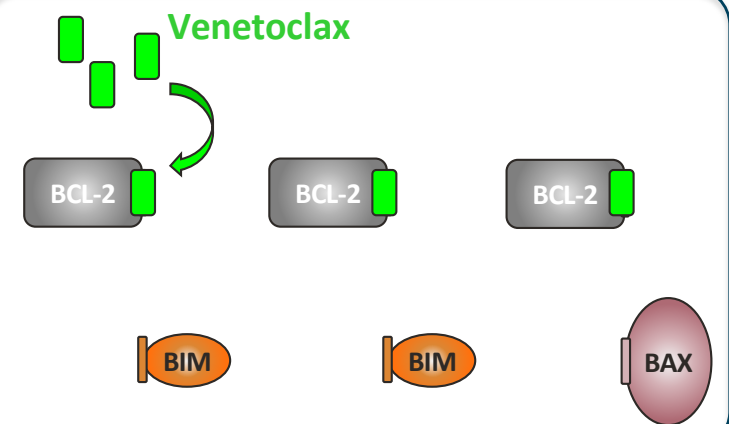
Pro-survival proteins (e.g. BCL-2) sequester pro-death proteins by binding to their **BH3 motifs**^{1,2}



Venetoclax³

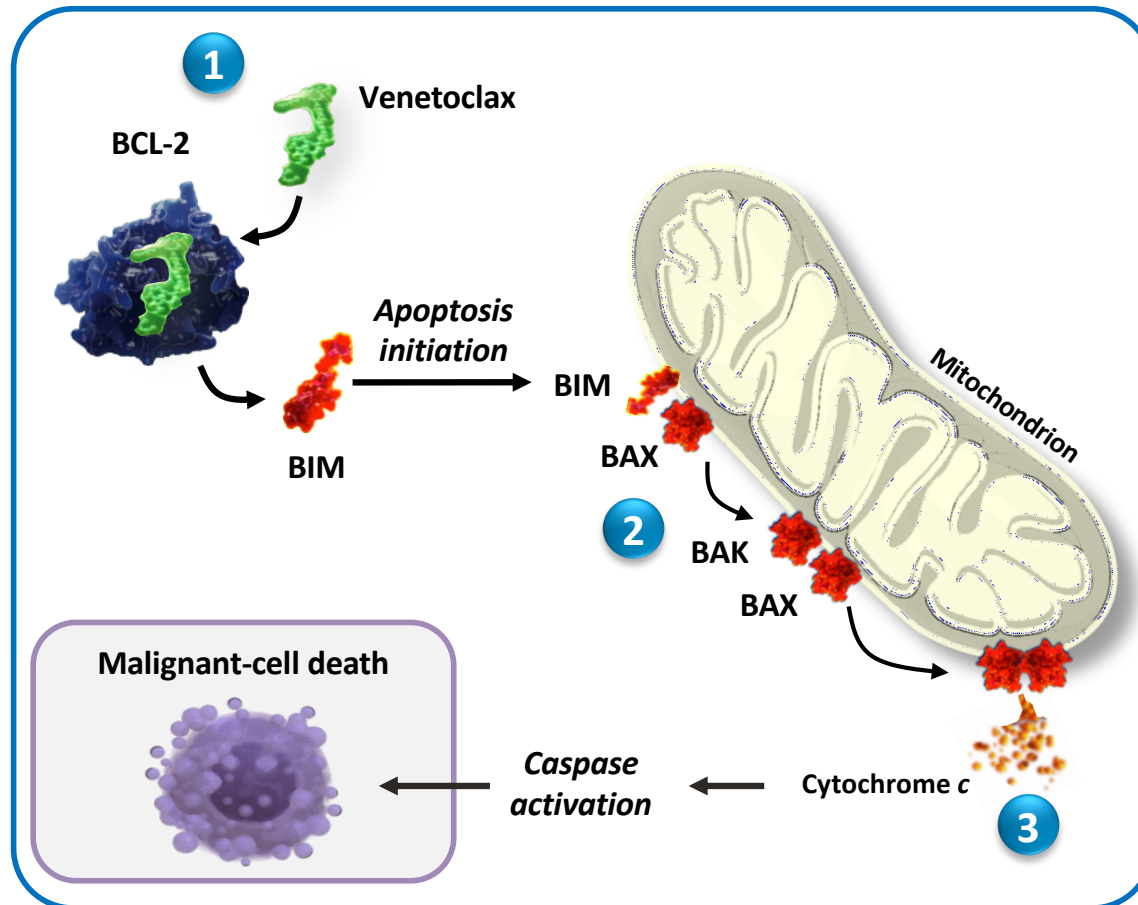


Venetoclax competes for binding to pro-survival proteins, freeing pro-death proteins³⁻⁵



1. Plati J, et al. *Integr Biol (Camb)* 2011; **3**:279–296; 2. Czabotar PE, et al. *Nat Rev Mol Cell Biol* 2014; **15**:49–63; 3. Souers AJ, et al. *Nat Med* 2013; **19**:202–208 (incl. suppl.); 4. Oltersdorf T, et al. *Nature* 2005; **435**:677–681; 5. Tse C, et al. *Cancer Res* 2008; **68**:3421–3428.

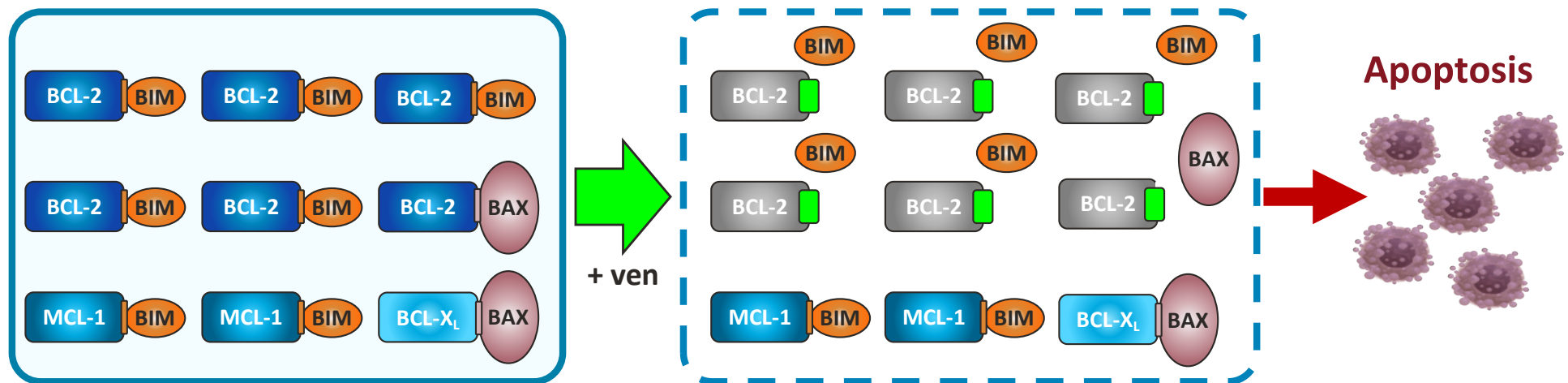
Venetoclax Induces Apoptosis in Malignant Cells



- 1** Venetoclax binds to BCL-2 and frees pro-death proteins (BIM, BAX)
- 2** The freed pro-death proteins interact and initiate an apoptotic cascade, which results in formation of a pore in the outer mitochondrial membrane
- 3** Cytochrome *c* is released from mitochondria and activates caspases, which dismantle the cell, resulting in malignant-cell death (apoptosis)

Malignant Cells with High Dependence on the BCL-2 Protein for Survival Are Inherently Sensitive to Venetoclax

Highly dependent on BCL-2 (increased levels of BCL-2 relative to other pro-survival proteins, including MCL-1 and BCL-X_L)



Inhibition of BCL-2 can free enough BIM and BAX to trigger apoptosis

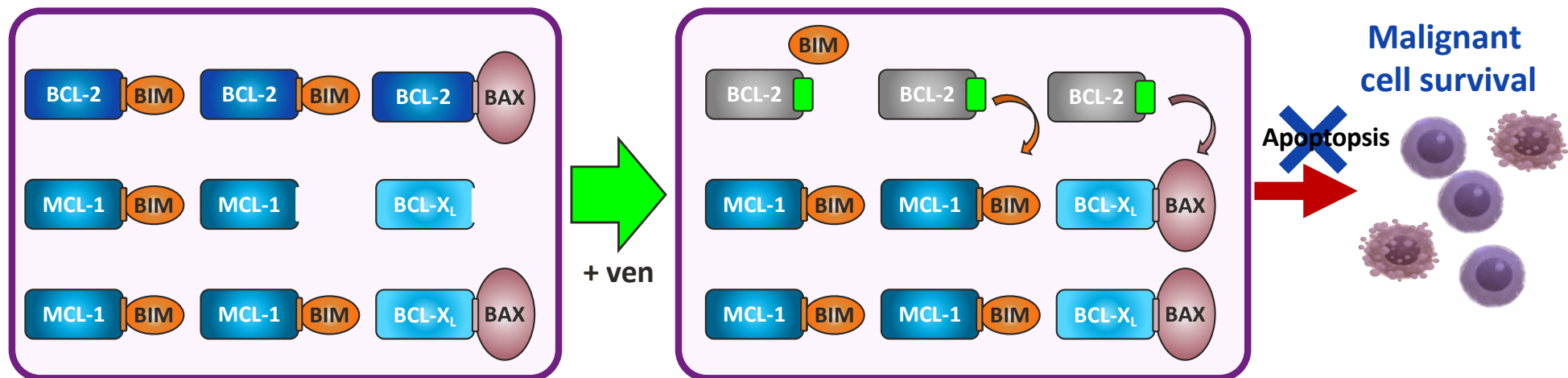
Key:

- Bound BIM–BCL-2
- Free BIM & inactivated BCL-2
- Venetoclax

1. Certo M, et al. *Cancer Cell* 2006; **9**:351–365;
2. Souers AJ, et al. *Nat Med* 2013; **19**:202–208 (incl. suppl.);
3. Punnoose EA, et al. *Mol Cancer Ther* 2016; **15**:1132–1144;
4. Levenson JD, et al. *Cancer Discov* 2017; **7**:1376–1393.




Malignant Cells That Are Co-dependent on Other Pro-survival Proteins May Be Less Sensitive to BCL-2 Inhibition Alone

Co-dependent on pro-survival proteins
(increased levels of other pro-survival proteins, MCL-1 and BCL-X_L)



Inhibition of BCL-2 may not free enough BIM and BAX to trigger apoptosis in all malignant cells

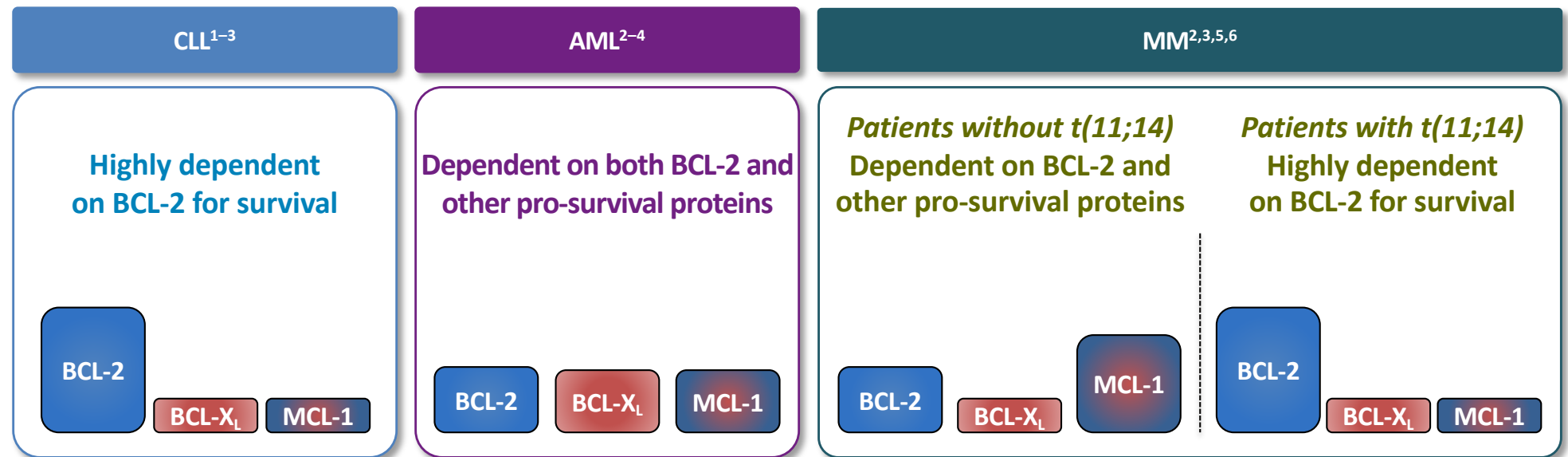
Key:

-  Bound BIM-BCL-2
-  Free BIM & inactivated BCL-2
-  Venetoclax

Patterns of expression of BCL2 family in selected lymphoid malignancies

| | Major | Level of bcl2 expression | Variability | Comment on mechanism |
|----------------------------|-------------------------|---|---------------------------------------|--|
| CLL | BCL2> MCL>> BCLxL | High | Some variability but always high | BCL2: loss of repression by miRNA15/16 MCL1 and BCLxL induced by CD40 ligation and microenviromental stimuli |
| Follicular Lymphoma | BCL2, MCL1, BCLxL | High | Rare to not be expressed | T(14;18) leads to constitutive expression. CD40L stimulates BCLxL expression MCL1 in centroblasts |
| DLBCL | BCL2> MCL1 | High GC type low in many ABC type | High where MYC driven MCL1>BCL2 | Varies including gene apmplification t(14;18) in double hit lymphoma, consequences of MYC dysregualtion |
| MCL | BCL2> MCL1 | High | Minor | Consequence of cyclin D1 dysregualtion MCL1 high in blastoid |
| Myeloma | BCL2, MCL1, BCLxL | High Especially t(11;14) | Moderate | BCL2 consequence of cyclin dysregulation MCL1 constitutively expressed in plasma cells BCLXL increased through microenviroment stimulation |
| ALL | BCL2, MCL1, BCLxL | variable | Significant | Appears to mimic expression pattern of precursor cell Patterned by oncogenic driver |

Malignant Cell Dependence on BCL-2 for Survival Has Been Shown across Several Hematologic Malignancies



Size of rectangles indicates relative dependency on specific protein for survival

Malignant cells' dependence on pro-survival proteins makes the BCL-2 family members rational targets for anticancer therapies

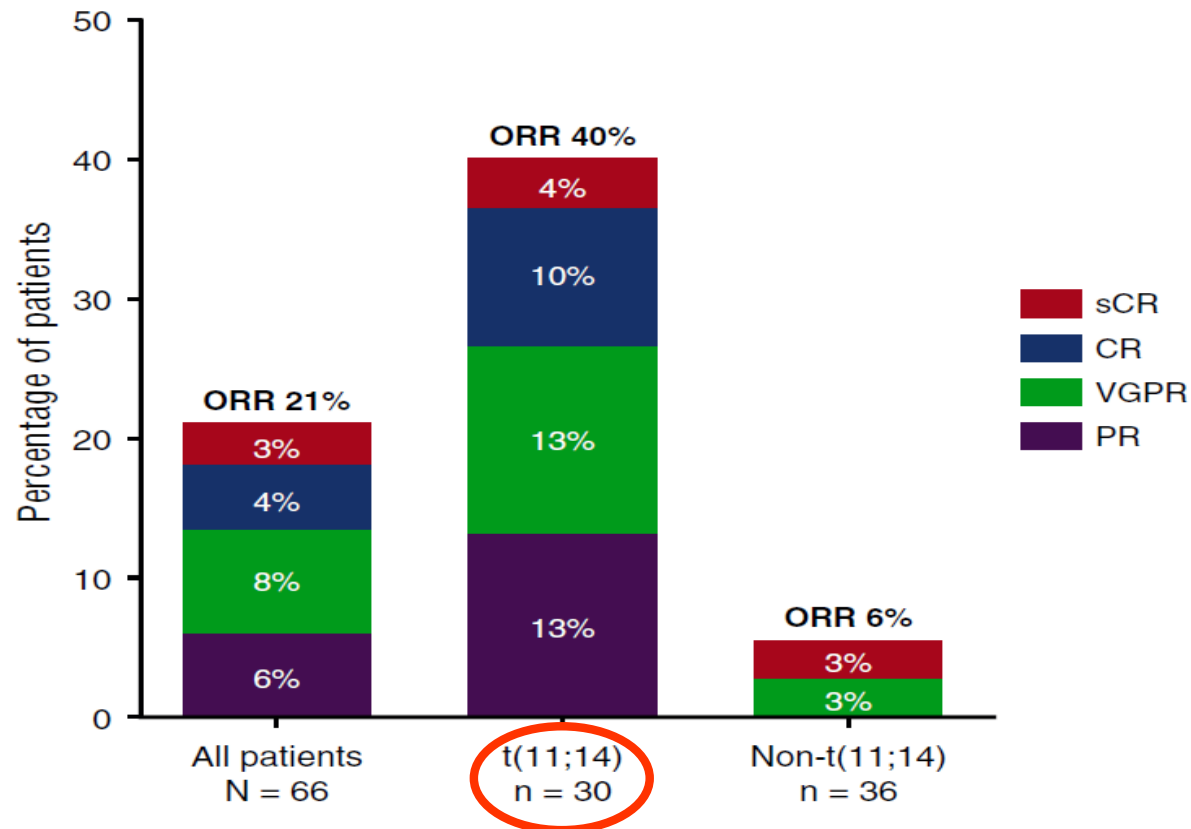
1. Del Gaizo Moore V, et al. *J Clin Invest* 2007; **117**:112–121; 2. Levenson JD, et al. *Cancer Discov* 2017; **7**:1376–1393;
3. Valentin R, et al. *Blood* 2018; **132**:1248–1264; 4. Pan R, et al. *Cancer Discov* 2014; **4**:362–375;
5. Touzeau C, et al. *Leukemia* 2018; **32**:1899–1907; 6. Gomez-Bougie P, et al. *Blood* 2018; **132**:2656–2669.

Rationale for anti-BCL2 agents use in multiple myeloma

- Gene-expression profiling (GEP) has shown that MM harboring t(11;14) has a specific pattern, characterized by CCND1 overexpression.
- GEP profile of t(11;14) plasma cells share similarities with GEP profile of lymphoma cells.
- Interestingly, t(11;14) plasma cells frequently express CD20, unlike all other myeloma subgroups.
- Plasma cells harboring t(11;14) are associated with increased dependency upon BCL2 for myeloma cell survival.

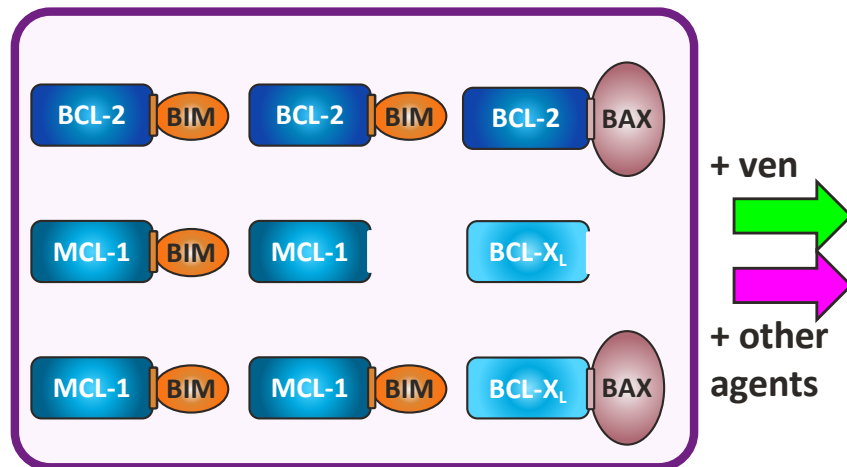
Venetoclax for t(11;14) MM

ORR rate by t(11;14) status



Venetoclax Can Combine with Agents That Increase BCL-2 Dependency or Target Other Pro-survival Proteins

Co-dependent on pro-survival proteins
(increased levels of other pro-survival proteins, MCL-1 and BCL-X_L)

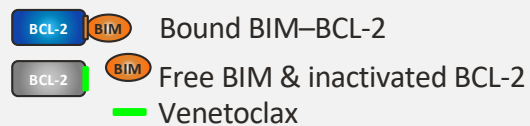


Sensitivity to BCL-2 inhibition can be promoted by increasing BCL-2 dependency (upregulation of BIM and BCL-2)*

and/or

Sensitivity to BCL-2 inhibition can be promoted by directly or indirectly downregulating or neutralizing other pro-survival proteins[†]

Key:

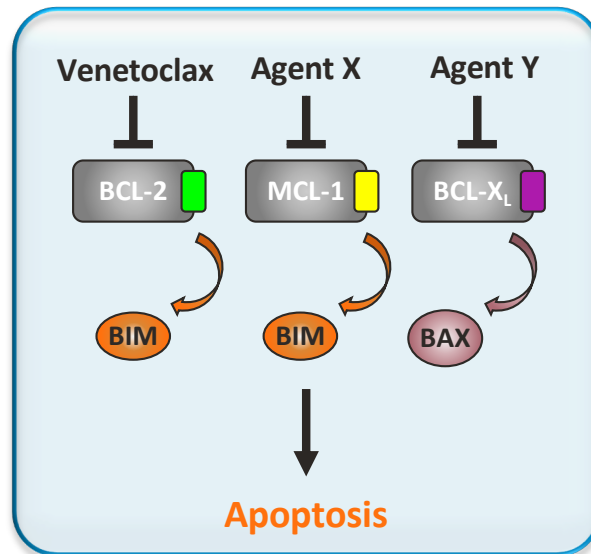


* e.g. dexamethasone; [†] e.g. proteasome inhibitors, HMAs, or LDAC.
HMA, hypomethylating agent; LDAC, low-dose cytarabine.

Venetoclax Can Also Combine with Agents Targeting Apoptotic and/or Non-apoptotic Complementary MoAs

Overlapping MoAs

Direct intrinsic apoptosis

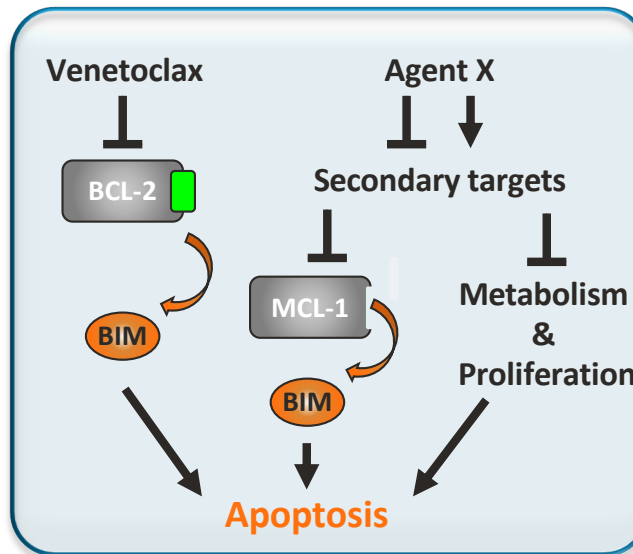


- MCL-1 inhibitors (S63845)
- BCL-X_L inhibitors (navitoclax)

Key: → Activation —| Inhibition

Partially Overlapping MoAs

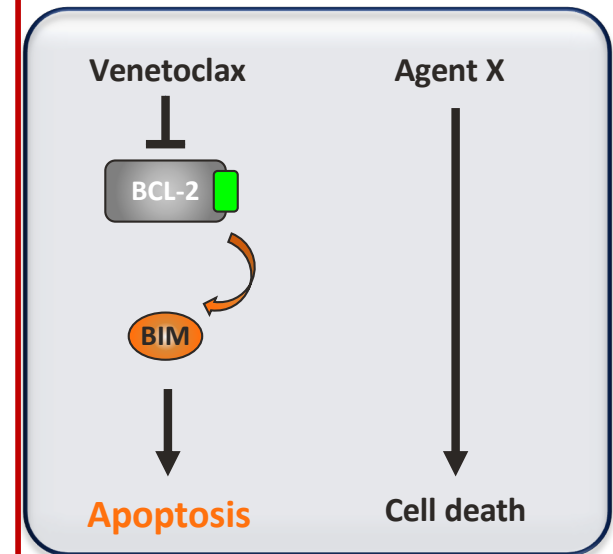
Indirect apoptosis + pleiotropic effect



- HMA, LDAC
- CDK9 inhibitors (alvocidib, dinaciclib)
- PIs (bortezomib, carfilzomib)
- FLT3 inhibitors (midostaurin, quizartinib, gilteritinib)
- MDM2 inhibitor (idasanutlin)






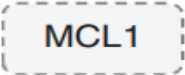
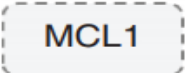
Distinct MoAs

Cell death other than apoptosis

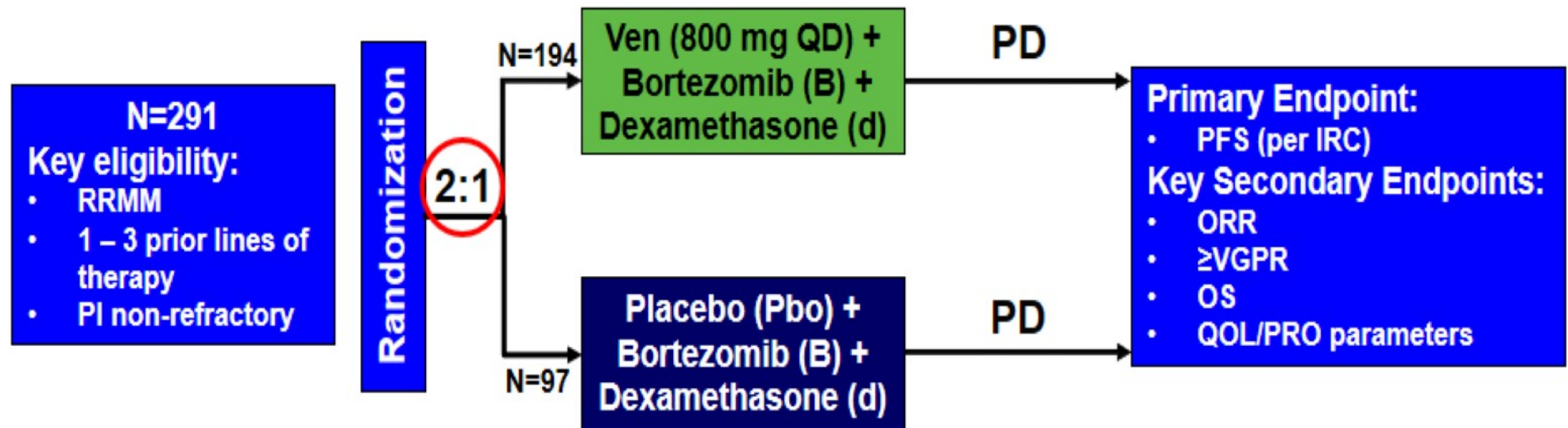


- Anti-CD20 antibodies (rituximab, obinutuzumab)
- Anti-CD38 antibodies (daratumumab)
- IMiDs (lenalidomide)
- Checkpoint inhibitors (pembrolizumab, nivolumab)

Rational combinations with the BCL2-selective inhibitor venetoclax

| | Mechanism | Agent(s) | Key target |
|---|--|---|------------------------|
|  | Direct inhibitor, BH3 mimetic | Venetoclax S55746 (BCL201) | BCL2 |
|  | Elevated BCL2 priming–BIM stabilization/upregulation | UO126, selumetinib Ibrutinib Dexamethasone Tamoxifen | MEK BTK HR ER |
|  | Direct inhibitor, BH3 mimetic | Navitoclax A-1331852 A-1155463 WEHI-539 | BCLX _L |
|  | Direct inhibitor, BH3 mimetic | S63845 A-1210477 | MCL1 |
|  | Reduced proteolysis of NOXA, a BH3-only protein which can bind/neutralize MCL1 | Bortezomib | Proteasome |
|  | Reduced RNA POLII-mediated transcriptional elongation of <i>MCL1</i> mRNA | Dinaciclib Alvocidib | CDK9 |
|  | Inhibition of cap-dependent translation of <i>MCL1</i> | AZD8055 | mTORC1/2 |
| | Unknown | Anthracyclines Hypomethylators | DNA/RNA DNA/RNA |

BELLINI Study Design



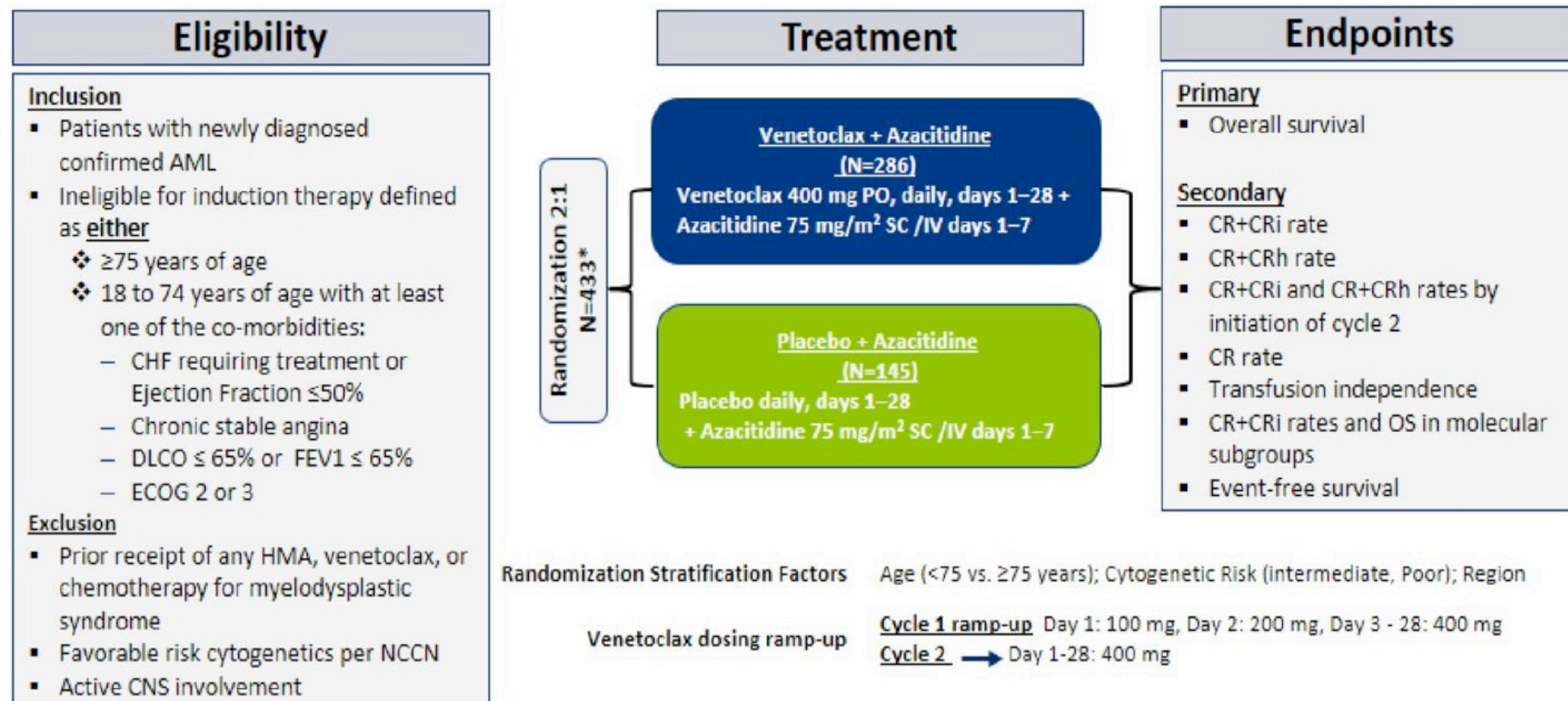
Cycles 1 – 8: 21-day, Bortezomib 1.3 mg/m² Days 1, 4, 8, 11 and dexamethasone 20 mg Days 1, 2, 4, 5, 8, 9, 11, 12

Cycles 9+: 35-day, Bortezomib 1.3 mg/m² Days 1, 8, 15, 22 and dexamethasone 20 mg Days 1, 2, 8, 9, 15, 16, 22, 23

| | |
|--------------------------------|---|
| Stratification factors | <ul style="list-style-type: none"> • Bortezomib sensitive vs naïve • Prior lines of therapy: 1 vs 2–3 |
| Non-ranked secondary endpoints | PFS in BCL-2 ^{high} (IHC), DOR, TTP, MRD negativity rate, other PROs (GHS, fatigue) |
| Key subgroup analyses | t(11;14), high/standard-risk cytogenetics, and BCL2 expression (gene expression) |

DOR, duration of response; GHS, global health status; IHC, immunohistochemistry; MRD, minimal residual disease; ORR, objective response rate; OS, overall survival; PFS, progression-free survival; PI, proteasome inhibitor; PRO, patient reported outcome; QD, daily; QOL, quality of life; RRMM, relapsed/refractory multiple myeloma; TTP, time to progression; VGPR, very good partial response.

VIALE-A Study Design (NCT02993523)



Take Home Messages

- Malignant cell dependence on pro-survival proteins makes the BCL-2 family members rational targets for anticancer therapies
- Malignant cells with high dependence on the BCL-2 protein for survival (like CLL cells) are Inherently sensitive to BCL-2 inhibition with Venetoclax
- However, malignant cells that are co-dependent on other pro-survival proteins (like AML and MM cells) may be less sensitive to BCL-2 inhibition alone.
- For this reason, Venetoclax can combine with agents targeting apoptotic and/or non-apoptotic complementary mechanism of action in diseases like AML, MM and potentially also NHL