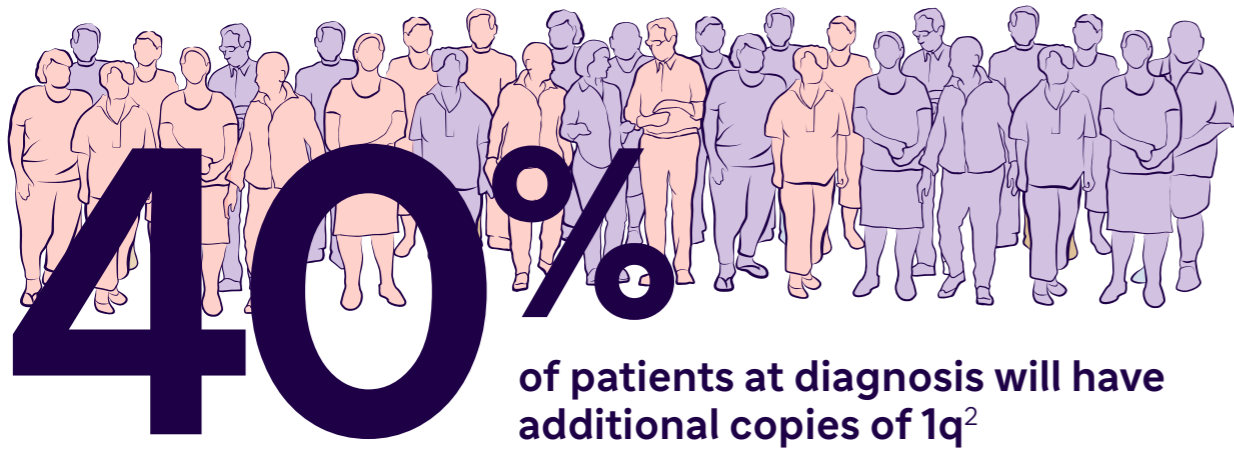


Multiple Myeloma: Chromosome 1q+ cytogenetic abnormalities

Multiple copies of chromosome 1 (1q+) is a frequent CA in MM

- MM is characterized by CAs in plasma cells¹
- Frequent gains include 1q, 6p and 11q¹
- 1q+ is one of the most common CAs in MM



Overexpression of certain genes in the 1q arm may drive pathogenesis and drug resistance^{4,5}

Key genes located in the 1q21 band:

- **BCL9**: Cell growth
- **ILF2, ADAR**: RNA metabolism
- **MCL1**: Anti-apoptosis
- **CKS1B**: Cell proliferation

- **PSMD4**: Anti-PI

Resistance to PIs through the upregulation of the proteasome genes, including PSMD4⁶

Other key genes outside of the 1q21 band include:

- **SLAMF7, SLAMF3**: Immunomodulatory receptors
- **FCRH5, NEK2**: Cell development
- **CD1D**: T-cell responses

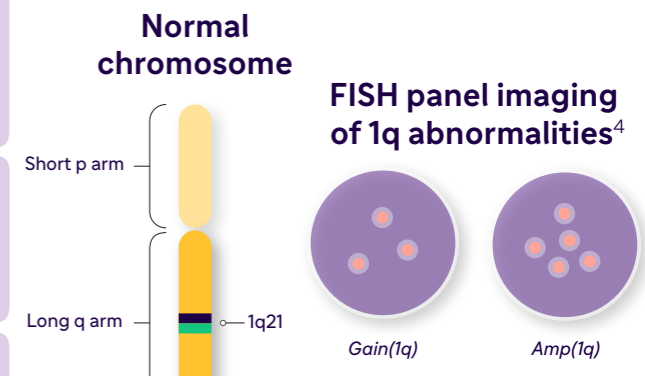
- **CD55, CD46**: CDC inhibition

Resistance to drugs that rely mainly on CDC through upregulation of complement inhibitors, such as CD55⁷

1q+ abnormalities are defined by copy number

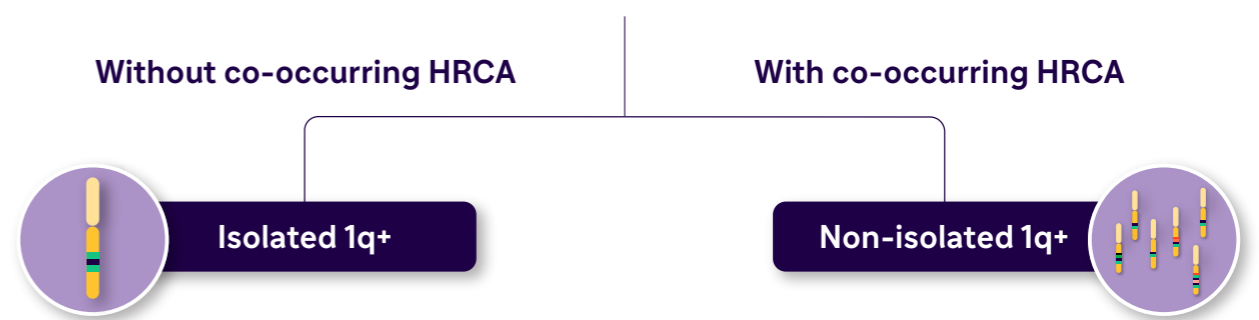
Chromosome 1q abnormalities terminology³

+1q	Additional copies of any part of the long arm of chromosome 1 (1q), irrespective of copy number or DNA segment gained
Gain(1q)	Gain of only 1 extra copy of chromosome 1q (3 total copies)
Amp(1q)	"Amplification" of 1q, with ≥ 2 additional copies of chromosome 1q (4 or more total copies)



In MM, 1q+ is frequently denoted as 1q21+, which refers to a specific location – region 2, band 1 on the long arm of chromosome 1

1q+ terminology when co-occurring with/without other high-risk cytogenetic abnormalities (HRCA)⁸



1q+ increases throughout the MM disease course^{4,9}

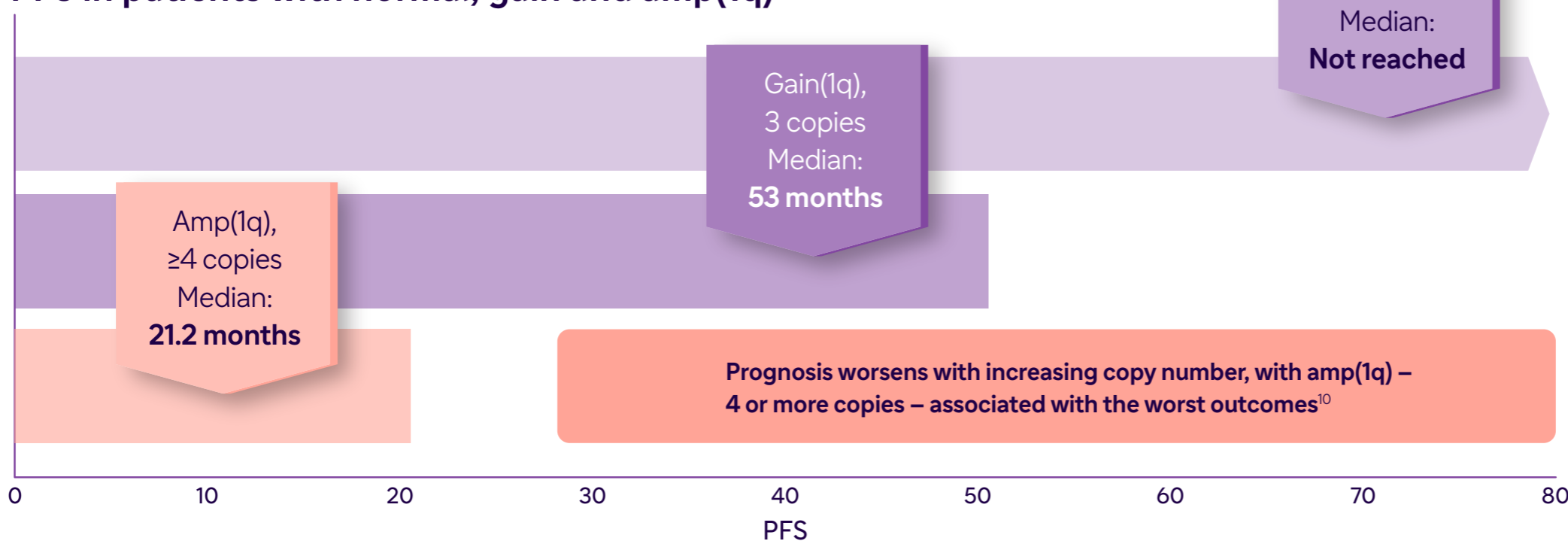
1q+: ~40% of NDMM cases⁴

1q+: 50–80% of RRMM cases⁴



1q+ is associated with poor outcomes, especially when presenting with other CAs

PFS in patients with normal, gain and amp(1q)



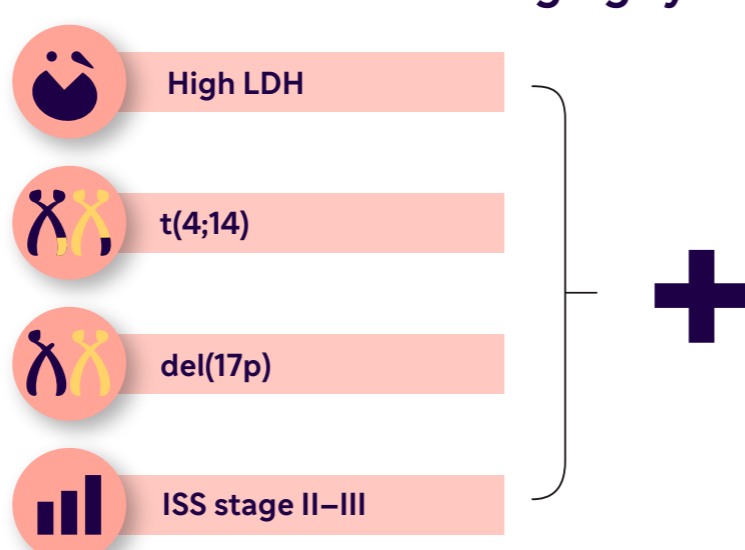
Prognosis worsens with increasing copy number, with amp(1q) – 4 or more copies – associated with the worst outcomes¹⁰

PFS and OS are significantly worse in patients with 1q+ versus patients without 1q abnormalities¹⁰

Co-occurrence of 1q+ with other high-risk factors has been associated with very poor prognosis (ultra-high-risk)^{11,5}

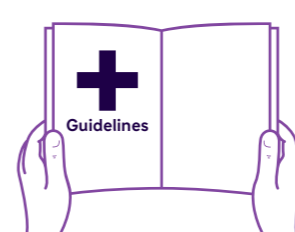
Incorporating 1q+ testing into clinical practice remains challenging

High-risk factors included in current staging systems^{12–14}



New staging systems, such as R2-ISS, MASS and R-ISS-1q,^{12–14} are including gain/amp(1q), alongside other high-risk factors when defining disease stage, in order to refine predictions of clinical outcomes and highlight the prognostic impact of gain/amp(1q)

Although most guidelines include 1q+ in their HRCA definitions, the recommendations for FISH panel testing for 1q+ are inconsistent and there is little treatment guidance specific to gain/amp(1q)^{15–17}



IMWG guidelines include gain/amp(1q) in their definition of HRCAs, but do not include it as part of their recommended routine FISH panel¹⁵

CA, cytogenetic abnormality; CDC, complement-dependent cytotoxicity; FISH, fluorescence in situ hybridization; HRCA, high-risk cytogenetic abnormality; IMWG, International Myeloma Working Group; ISS, international staging system; LDH, lactate dehydrogenase; MASS, The Mayo Additive Staging System; MM, multiple myeloma; MRD, minimal residual disease; NDMM, newly-diagnosed multiple myeloma; OS, overall survival; PFS, progression-free survival; PI, protease inhibitor; R-ISS, Revised International Staging System; RRMM, relapsed/refractory multiple myeloma.

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