

## PLASMA CELL NEOPLASMS (PCN)

### DISEASE

**Note: complete this form only if this diagnosis was the indication for the HCT/CT or if it was specifically requested. Consult the manual for further information.**

**Date of diagnosis:** \_\_\_\_/\_\_\_\_/\_\_\_\_ (YYYY/MM/DD)

**Classification (WHO 2022):**

<input type="checkbox"/> Plasma cell (multiple) myeloma (PCM)	<input type="checkbox"/> Heavy chain and light chain  <input type="checkbox"/> Light chain only	<b>Heavy chain type:</b> <input type="checkbox"/> IgG <input type="checkbox"/> IgA <input type="checkbox"/> IgD <input type="checkbox"/> IgE <input type="checkbox"/> IgM (not Waldenstrom) <input type="checkbox"/> Unknown	<b>Light chain type:</b> <input type="checkbox"/> Kappa <input type="checkbox"/> Lambda <input type="checkbox"/> Unknown
<input type="checkbox"/> Non-secretory			
<input type="checkbox"/> Unknown			
<input type="checkbox"/> Plasma cell leukaemia			
<input type="checkbox"/> Solitary plasmacytoma of bone			
<input type="checkbox"/> Immunoglobulin-related (AL) amyloidosis			
<input type="checkbox"/> POEMS (Polyneuropathy, Organomegaly, Endocrinopathy/Edema, Monoclonal-protein, Skin changes)			
<input type="checkbox"/> Monoclonal immunoglobulin deposition disease			
<input type="checkbox"/> Other; specify: _____			

## STAGING

*PCM only*

**Staging at diagnosis:**
**Revised ISS:**

Stage
<input type="checkbox"/> I: ISS I without high risk FISH (del(17p) and/or t(4;14) and/or t(14;16) and normal LDH
<input type="checkbox"/> II: not R-ISS I or III
<input type="checkbox"/> III: ISS III with high risk FISH (del(17p) and/or t(4;14) and/or t(14;16)) and/or high LDH
<input type="checkbox"/> Unknown

**ISS:**

Stage	β2-μglob (mg/L)	Albumin (g/L)
<input type="checkbox"/> I	< 3.5	> 35
<input type="checkbox"/> II	< 3.5 OR 3.5 ≤ 5.5	< 35 any
<input type="checkbox"/> III	> 5.5	any
<input type="checkbox"/> Unknown		

**Extramedullary disease (EMD):**

<input type="checkbox"/> No					
<input type="checkbox"/> Yes	EMD diagnosed on MRI	<input type="checkbox"/> No	<input type="checkbox"/> Yes	<input type="checkbox"/> Unknown	
	EMD diagnosed on PET-CT	<input type="checkbox"/> No	<input type="checkbox"/> Yes	<input type="checkbox"/> Unknown	
	Location of EMD	<input type="checkbox"/> Paraskeletal	<input type="checkbox"/> Organ	<input type="checkbox"/> Both	<input type="checkbox"/> Unknown
	Specify organ: _____				
<input type="checkbox"/> Unknown					



EBMT Centre Identification Code (CIC): \_\_\_\_\_

Treatment Type  HCT  CT  GT  IST  Other

Hospital Unique Patient Number (UPN): \_\_\_\_\_

Patient Number in EBMT Registry: \_\_\_\_\_

Treatment Date \_\_\_\_/\_\_\_\_/\_\_\_\_ (YYYY/MM/DD)

**CHROMOSOME ANALYSIS****Chromosome analysis done at diagnosis:**

- No  
 Yes:      **Output of analysis:**  Separate abnormalities       Full karyotype  
 Unknown

*If chromosome analysis was done:***What were the results?**

- Normal  
 Abnormal: number of abnormalities present: \_\_\_\_\_  
 Failed

Date of chromosome analysis: \_\_\_\_/\_\_\_\_/\_\_\_\_ (YYYY/MM/DD)       UnknownChromosome analysis method used:  Karyotyping  
 FISH

Indicate below whether the abnormalities were absent, present or not evaluated.

<b>1q amplification (4 or more copies)</b>	<input type="checkbox"/> Absent	<input type="checkbox"/> Present	<input type="checkbox"/> Not evaluated	<input type="checkbox"/> Unknown
<b>1q gain (3 copies)</b>	<input type="checkbox"/> Absent	<input type="checkbox"/> Present	<input type="checkbox"/> Not evaluated	<input type="checkbox"/> Unknown
<b>abn(17q)</b>	<input type="checkbox"/> Absent	<input type="checkbox"/> Present	<input type="checkbox"/> Not evaluated	<input type="checkbox"/> Unknown
<b>del1p</b>	<input type="checkbox"/> Absent	<input type="checkbox"/> Present	<input type="checkbox"/> Not evaluated	<input type="checkbox"/> Unknown
<b>del(17p)</b>	<input type="checkbox"/> Absent	<input type="checkbox"/> Present	<input type="checkbox"/> Not evaluated	<input type="checkbox"/> Unknown
<b>del(13q14)</b>	<input type="checkbox"/> Absent	<input type="checkbox"/> Present	<input type="checkbox"/> Not evaluated	<input type="checkbox"/> Unknown
<b>Hyperdiploidy</b>	<input type="checkbox"/> Absent	<input type="checkbox"/> Present	<input type="checkbox"/> Not evaluated	<input type="checkbox"/> Unknown
<b>myc rearrangement</b>	<input type="checkbox"/> Absent	<input type="checkbox"/> Present	<input type="checkbox"/> Not evaluated	<input type="checkbox"/> Unknown
<b>t(4;14)</b>	<input type="checkbox"/> Absent	<input type="checkbox"/> Present	<input type="checkbox"/> Not evaluated	<input type="checkbox"/> Unknown
<b>t(6;14)</b>	<input type="checkbox"/> Absent	<input type="checkbox"/> Present	<input type="checkbox"/> Not evaluated	<input type="checkbox"/> Unknown
<b>t(11;14)</b>	<input type="checkbox"/> Absent	<input type="checkbox"/> Present	<input type="checkbox"/> Not evaluated	<input type="checkbox"/> Unknown
<b>t(14;16)</b>	<input type="checkbox"/> Absent	<input type="checkbox"/> Present	<input type="checkbox"/> Not evaluated	<input type="checkbox"/> Unknown
<b>t(14;20)</b>	<input type="checkbox"/> Absent	<input type="checkbox"/> Present	<input type="checkbox"/> Not evaluated	<input type="checkbox"/> Unknown
Other; specify: _____	<input type="checkbox"/> Absent	<input type="checkbox"/> Present		

OR

Transcribe the complete karyotype: \_\_\_\_\_