KU LEUVEN



RegaVir platform: Case discussions antiviral resistance testing

Robert Snoeck & Graciela Andrei

Leuven, April 26, 2022



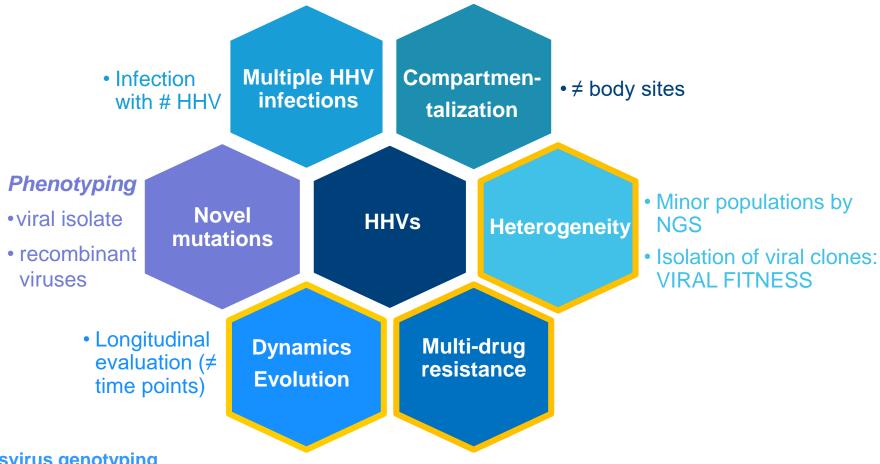
Cytomegalovirus (CMV) infection in hematopoietic stem cell transplantation (HSCT) patients

- Prophylactic or preemptive therapy against CMV infection has successfully reduced the incidence of CMV disease in transplant recipients.
- However, CMV still remains one of the major infectious complications and causes of morbidity and mortality among HSCT recipients.
- Risk factors for CMV infection in HSCT patients:
 - CMV seropositivity of donor and recipient.
 - Allogeneic transplantation (especially with T cell-depleted unrelated or HLA-mismatched donors).
 - Cell source: cord blood recipients have both longer and higher cumulative incidences of CMV infection compared to peripheral blood or allogeneic bone marrow recipients from related or unrelated donor origin.



Cytomegalovirus (CMV) infection in hematopoietic stem cell transplantation (HSCT) patients

- In HSCT patients, cellular immunity is mainly significantly impaired during the **first 100 days post-transplant**.
- CMV infection in HSCT patients manifests usually as interstitial pneumonia and gastrointestinal disease.
- Although less frequently, late CMV reactivation (>100 days posttransplant) can occur.
- A few reports have highlighted the **importance of CMV resistance in pediatric HSCT populations**.
 - GCV-R associated with the UL97 protein kinase (Ohta et al., 2001, Gohring et al., 2009, Kim et al., 2012, Erice et al., 1989).
 - ▶ GCV-R & PFA-R conferred by multiple UL97 and UL54 (DNA polymerase) mutations (Choi et al., 2014)
 - ➢ GCV-R, PFA-R & CDV-R conferred by DNA polymerase mutations (<u>Blackman et al., 2004</u>, <u>Drouot et al., 2014</u>, <u>Springer et al., 2005</u>).



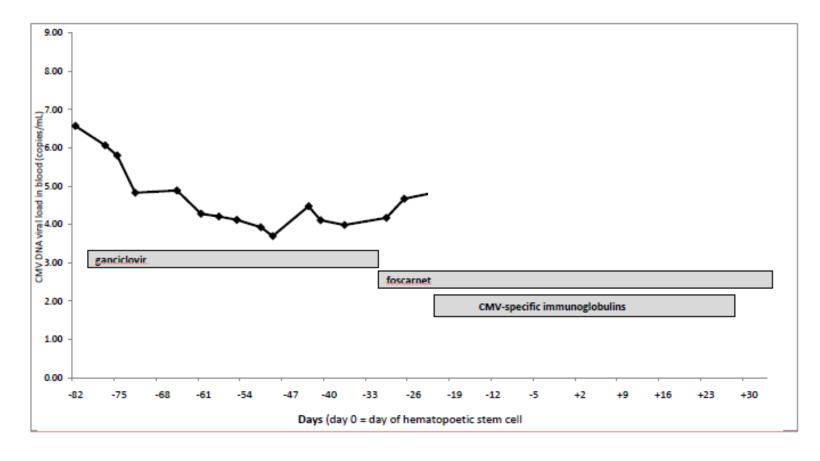
Herpesvirus genotyping

- **Prospectively:** capillary (Sanger sequencing)
- Retrospectively: next-generation sequencing (NGS) •



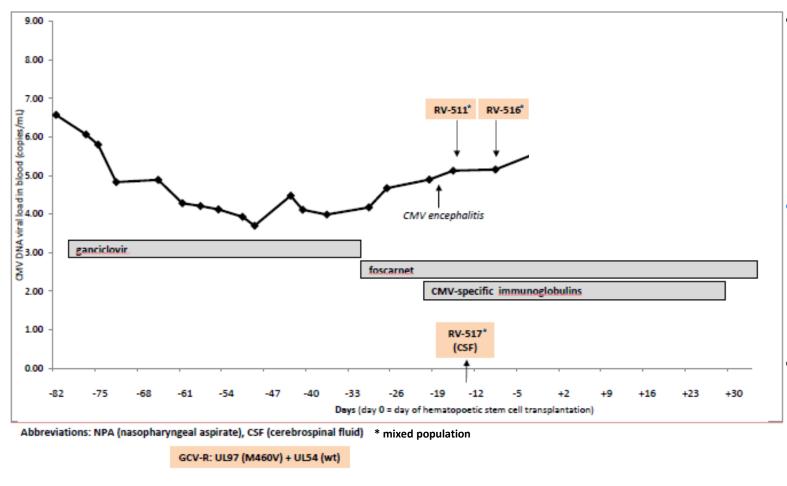
- A 4-month-old girl (weight 5.8 kg) was referred for fever, failure to thrive and bloody diarrhea.
- She was the first child of **consanguineous parents**, so **severe immunodeficiency** was presumed.
- CMV was detected in the blood: viral load of 3.77 10E6 copies/ml \rightarrow GCV (5 mg/kg, 2x/d) started.
- MHC class II deficiency was diagnosed and the search for a stem cell donor was initiated.



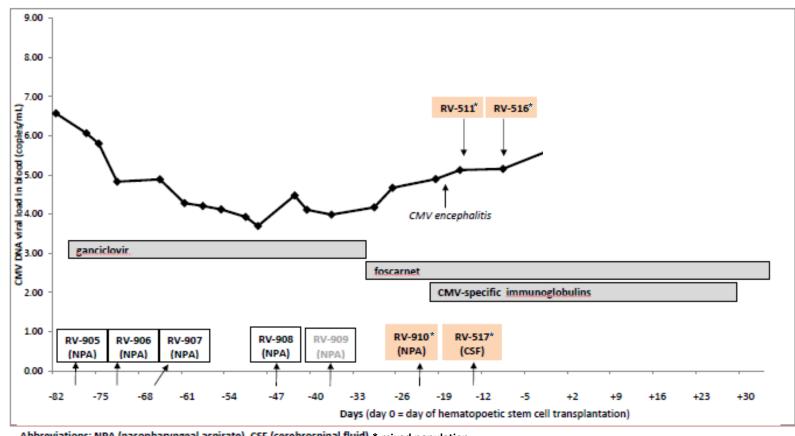


- Four weeks later the dose frequency of GCV was reduced to 1x/d.
- However, 1 week later, the CMV viral load started to increase → GCV was switched from 1x/d to 2x/d.
- As the CMV viral load still increased, GCV administration was stopped, and therapy was switched to PFA.
- No HLA identical donor nor cord blood was found so haploidentical HSCT was planned.
- For more than one month, the patient received PFA (2 × 350 mg) and simultaneously CMV-specific immunoglobulins.





- Antiviral resistance testing in a **blood** sample **(RV-511)** was performed and revealed the presence of a mixed population of **M460V-mutant** and wild-type virus in the UL97 protein kinase.
- M460V UL97 mutation is known to confer GCV-R with susceptibility to foscarnet and cidofovir (Lurain and Chou, 2010).
 - Despite antiviral treatment the patient deteriorated and developed CMV encephalitis confirmed by positivity of CMV PCR on CSF taken on day –14 (RV-517), which showed a similar resistance profile as the blood samples (RV-511 and RV-516)



- Retrospective analysis of NPA samples (RV-905 to RV-910) was performed.
- The NPA sample taken during the same time period (RV-910) had a mixed population of M460V-mutant and wild-type virus in the UL97 protein kinase
- NPA samples recovered at earlier time points (RV-905 to RV-908) showed a wild-type CMV genotype, indicating that GCV-R emerged at the time the patient developed encephalitis.

Abbreviations: NPA (nasopharyngeal aspirate), CSF (cerebrospinal fluid) * mixed population

GCV-R: UL97 (M460V) + UL54 (wt)



Specimens recovered before transplantation (day 0 = day of HSCT)

Retrospective analysis)

Prospective analysis)

| | Day | Samples code | CMV DNA | Type of sample | Amino acids changes related to resistance in: | | Resistance to: | Amino acid changes related to genetic polymorphisms | |
|--|-----|---------------|-----------------|----------------|---|----------------|----------------|---|------------------------------------|
| | | | | | UL97 (protein kinase) | UL54 (DNA pol) | | UL97 (protein kinase) | UL54 (DNA polymerase) |
| | -82 | Not available | 3.77 × 10e6 | Blood | Not available | Not available | Not available | Not available | Not available |
| | -79 | RV-905 | Positive | NPA | None | None | None | None | S655L, F669L, N685S, A885T & N898D |
| | -72 | RV-906 | Positive | NPA | None | None | None | None | S655L, F669L, N685S, A885T & N898D |
| | -65 | RV-907 | Positive | NPA | None | None | None | None | S655L, F669L, N685S, A885T & N898D |
| | -43 | RV-908 | Positive | NPA | None | None | None | None | S655L, F669L, N685S, A885T & N898D |
| | -37 | RV-909 | Detection limit | NPA | Not available | Not available | _ | Not available | Not available |
| | -19 | RV-910 | Positive | NPA | M460V* | None | GCV | None | S655L, F669L, N685S, A885T & N898D |
| | -16 | RV-511 | 1.34 × 10e5 | Blood | M460V* | none | GCV | None | S655L, F669L, N685S, A885T & N898D |
| | -14 | RV-517 | Positive | CSF | M460V* | none | GCV | None | S655L, F669L, N685S, A885T & N898D |
| | -9 | RV-516 | 1.44 × 10e5 | Blood | M460V* | none | GCV | None | S655L, F669L, N685S, A885T & N898D |

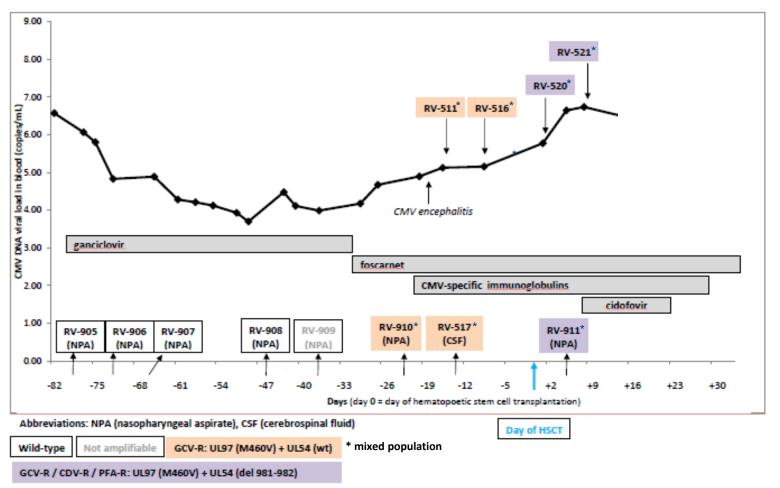
NA (not amplifiable), NPA (nasopharyngeal aspirate), CSF (cerebrospinal fluid), GCV (ganciclovir), CDV (cidofovir), FOS (foscarnet), del (deletion), DL (detection limit), ND (not detected).

Samples marked in bold were analyzed when they were recovered from the patient in order to adapt antiviral treatment while those indicated in italic were evaluated retrospectively.



- Despite CMV infection, an allogeneic transplantation was performed (day 0 = day of HSCT) with peripheral blood stem cells from a haploidentical donor, after selection of CD34 cells from the graft.
- Both the donor and receptor were CMV IgG seropositive (D+/R+).
- The **conditioning regimen** consisted of intravenous busulfan ($16 \times 1 \text{mg/kg}$) and fludarabine ($4 \times 35 \text{mg/m2}$).
- Three days after HSCT, the patient developed fever for which antibiotics were started.
- Because of progression of neurologic encephalitic disturbances suggestive of central CMV infection, prophylactic anti-epileptics were started (phenobarbital) and cidofovir was added empirically to the foscarnet antiviral treatment.





- Antiviral resistance testing of a blood sample on day +1 (RV-520):
 - mixed population of M460V-mutant and wild-type virus at the UL97 protein kinase.
 - mixed population of 981-982 deletion mutant and wild-type virus in the DNA polymerase.
- DNA pol 981-982 del. is known to confer resistance to ganciclovir, foscarnet and cidofovir (Lurain and Chou, 2010).
- The same viral populations were detected at the same time in the NPA (RV-911) on day +8 in blood. (RV-521).

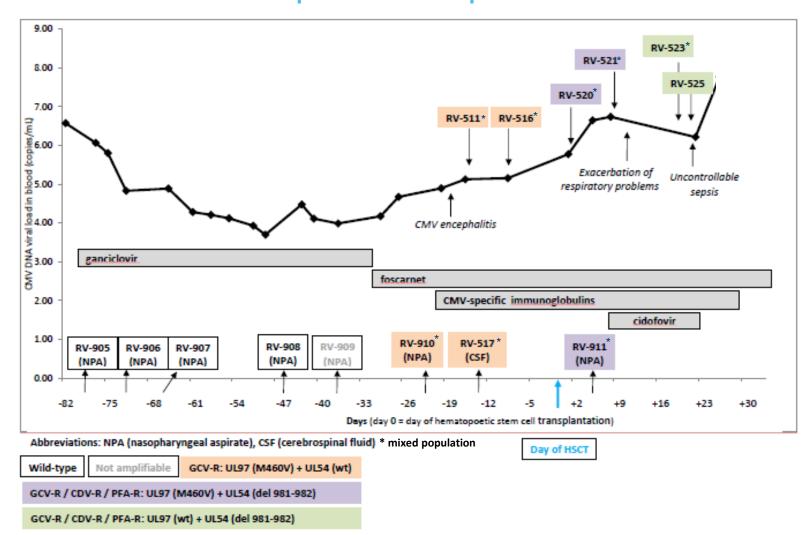


Specimens recoverted after day 0 (the day of transplantation)

| Day | Samples code | CMV DNA (copies/ml) | Type of sample | Amino acids changes related to resistance in: | | Resistance to: | Amino acid changes related to genetic polymorphisms | |
|-----|--------------|------------------------|----------------|---|----------------|----------------|---|------------------------------------|
| | | | | UL97 (protein kinase) | UL54 (DNA pol) | | UL97 (protein kinase) | UL54 (DNA polymerase) |
| +1 | RV-520 | 5.98 × 10e5 | Blood | M460V* | 981-982 del.* | GCV, CDV, FOS | None | S655L, F669L, N685S, A885T & N898D |
| +5 | RV-911 | Positive | NPA | M460V* | 981-982 del. * | GCV, CDV, FOS | None | S655L, F669L, N685S, A885T & N898D |
| +8 | RV-521 | 5.52 × 10e6 | Blood | M460V* | 981-982 del. * | GCV, CDV, FOS | N68D, I244V | S655L, F669L, N685S, A885T & N898D |
| | | | | | | | | |
| | | | | | | | | |
| | | | | | | | | |
| | | | | | | | | |
| | | | | | | | | |
| | | | | | | | | |
| | | | | | | | | |
| | | | | | | | | |

NA (not amplifiable), NPA (nasopharyngeal aspirate), CSF (cerebrospinal fluid), GCV (ganciclovir), CDV (cidofovir), FOS (foscarnet), del (deletion), DL (detection limit), ND (not detected).

Samples marked in bold were analyzed when they were recovered from the patient in order to adapt antiviral treatment while those indicated in italic were evaluated retrospectively.



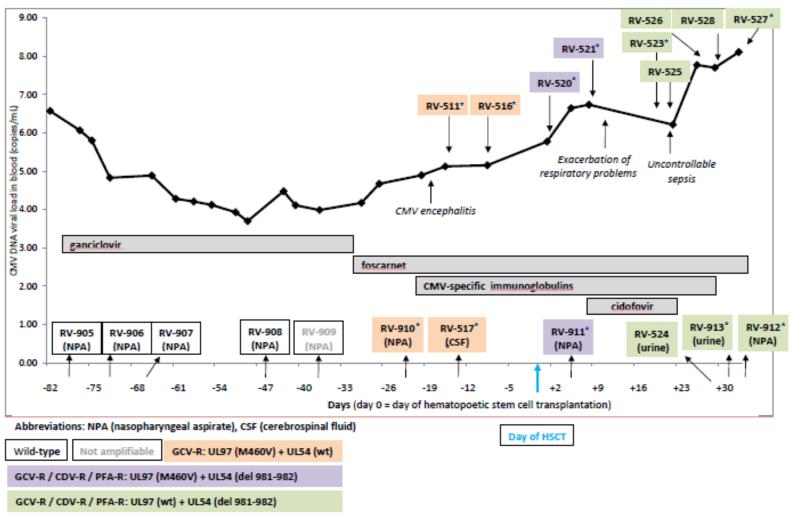
- Day +13, the patient was referred to the intensive care unit because of exacerbation of respiratory problems.
- Cidofovir was stopped because of severe cytopenia and increase of the CMV viral load.
- Day +19, in the blood sample (RV-523)
 - > UL97 M460V-mutant population was not identified anymore.
 - A mixed population of DNA polymerase mutant virus bearing the 981-982 deletion was present.
- Day +22, in only 3 days, the mutant virus totally replaced the wild-type virus (RV-525) and presented a pure population of DNA polymerase mutant virus with the 981-982 deletion.

Specimens recoverted after day 0 (the day of transplantation)

| Day | Samples code | CMV DNA (copies/ml) | Type of sample | Amino acids changes related to resistance in: | | Resistance to: | Amino acid changes related to genetic polymorphisms | |
|-----|--------------|------------------------|----------------|---|----------------|----------------|---|------------------------------------|
| | | | | UL97 (protein kinase) | UL54 (DNA pol) | | UL97 (protein kinase) | UL54 (DNA polymerase) |
| +1 | RV-520 | 5.98 × 10e5 | Blood | M460V* | 981-982 del.* | GCV, CDV, FOS | None | S655L, F669L, N685S, A885T & N898D |
| +5 | RV-911 | Positive | NPA | M460V* | 981-982 del. * | GCV, CDV, FOS | None | S655L, F669L, N685S, A885T & N898D |
| +8 | RV-521 | 5.52 × 10e6 | Blood | M460V* | 981-982 del. * | GCV, CDV, FOS | N68D, I244V | S655L, F669L, N685S, A885T & N898D |
| +19 | RV-523 | 1.06 × 10e7 | Blood | none | 981-982 del. * | GCV, CDV, FOS | None | S655L, F669L, N685S, A885T & N898D |
| +22 | RV-525 | 1.65 × 10e6 | Blood | none | 981-982 del. | GCV, CDV, FOS | None | S655L, F669L, N685S, A885T & N898D |
| | | | | | | | | |
| | | | | | | | | |
| | | | | | | | | |
| | | | | | | | | |
| | | | | | | | | |
| | | | | | | | | |

NA (not amplifiable), NPA (nasopharyngeal aspirate), CSF (cerebrospinal fluid), GCV (ganciclovir), CDV (cidofovir), FOS (foscarnet), del (deletion), DL (detection limit), ND (not detected).

Samples marked in bold were analyzed when they were recovered from the patient in order to adapt antiviral treatment while those indicated in italic were evaluated retrospectively.



- During the following days there was a further clinical deterioration of respiratory and liver function and uncontrollable sepsis.
- No access to compassionate use of novel anti-CMV drugs under clinical development.
- It was decided to stop antiviral treatment and to start palliative care.
- Multidrug-resistant CMV infection [due to the presence of the 981-982 deletion in the viral DNA polymerase not only in the blood (RV-526, RV-527, and RV-528) but also in urine (RV-524 and RV-913) and NPA (RV-912)] together with encephalitis, were all cumulating in a respiratory arrest with cardiac arrest and death of the patient at day +36.

Specimens recoverted after day 0 (the day of transplantation)

| Day | Samples code | CMV DNA (copies/ml) | Type of sample | Amino acids changes related to resistance in: | | Resistance to: | Amino acid changes related to genetic polymorphisms | |
|-----|--------------|------------------------|----------------|---|----------------|----------------|---|------------------------------------|
| | | | | UL97 (protein kinase) | UL54 (DNA pol) | | UL97 (protein kinase) | UL54 (DNA polymerase) |
| +1 | RV-520 | 5.98 × 10e5 | Blood | M460V* | 981-982 del.* | GCV, CDV, FOS | None | S655L, F669L, N685S, A885T & N898D |
| +5 | RV-911 | Positive | NPA | M460V* | 981-982 del. * | GCV, CDV, FOS | None | S655L, F669L, N685S, A885T & N898D |
| +8 | RV-521 | 5.52 × 10e6 | Blood | M460V* | 981-982 del. * | GCV, CDV, FOS | N68D, I244V | S655L, F669L, N685S, A885T & N898D |
| +19 | RV-523 | 1.06 × 10e7 | Blood | none | 981-982 del. * | GCV, CDV, FOS | None | S655L, F669L, N685S, A885T & N898D |
| +22 | RV-525 | 1.65 × 10e6 | Blood | none | 981-982 del. | GCV, CDV, FOS | None | S655L, F669L, N685S, A885T & N898D |
| +26 | RV-526 | 5.94 × 10e7 | Blood | none | 981-982 del. | GCV, CDV, FOS | None | S655L, F669L, N685S, A885T & N898D |
| +28 | RV-524 | Positive | Urine | none | 981-982 del. | GCV, CDV, FOS | None | S655L, F669L, N685S, A885T & N898D |
| +29 | RV-528 | 5.09 × 10e7 | Blood | none | 981-982 del. | GCV, CDV, FOS | None | S655L, F669L, N685S, A885T & N898D |
| +30 | RV-913 | Positive | Urine | none | 981-982 del. * | GCV, CDV,FOS | None | S655L, F669L, N685S, A885T & N898D |
| +31 | RV-912 | ND | NPA | none | 981-982 del. * | GCV, CDV, FOS | None | S655L, F669L, N685S, A885T & N898D |
| +33 | RV-527 | 1.29 × 10e8 | Blood | none | 981-982 del. * | GCV, CDV, FOS | Non | S655L, F669L, N685S, A885T & N898D |

NA (not amplifiable), NPA (nasopharyngeal aspirate), CSF (cerebrospinal fluid), GCV (ganciclovir), CDV (cidofovir), FOS (foscarnet), del (deletion), DL (detection limit), ND (not detected).

Samples marked in bold were analyzed when they were recovered from the patient in order to adapt antiviral treatment while those indicated in italic were evaluated retrospectively.

CMV disease of the CNS

- Has occurred mostly in patients with advanced AIDS, rarely reported in HSCT patients.
- Is a rare complication after HSCT or solid-organ transplantation.
- In our patient, CMV encephalitis was associated with:
 - recurrent CMV viremia following prolonged administration of ganciclovir and foscarnet.
 - emergence of drug-resistance.



CMV central nervous system (CNS) disease after allogeneic HSCT

• Is a late-onset disease (median time of onset, 210 days) and is usually manifested as encephalitis (Reddy et al, 2010)

Associated with:

- **T-cell depletion, anti-thymocyte globulin, umbilical cord blood transplantation** that cause severe and extended T-cell immunodeficiency (8 of 11 cases)
- a history of **recurrent CMV viremia** treated with multiple courses of preemptive ganciclovir or foscarnet therapy (11 of 11 cases)
- drug-resistance (11 of 11 cases)
- high mortality despite therapy with a combination of anti-CMV drugs (ganciclovir, foscarnet and cidofovir) (10 of 11 cases)



CMV central nervous system (CNS) disease after allogeneic HSCT

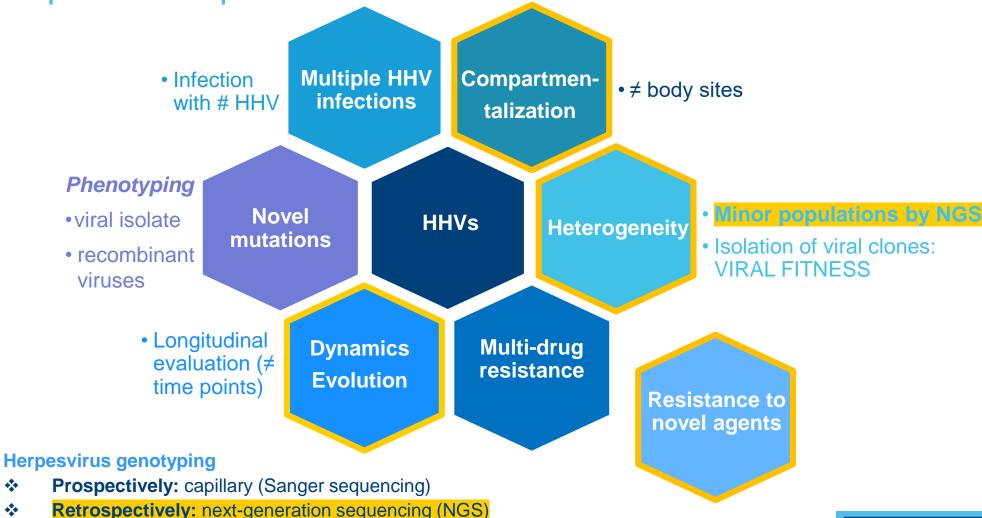
- > Suboptimal ganciclovir dose may promote the emergence of drug-resistance since adequate drug levels are crucial to control viral replication.
- ➢ High drug levels are known to delay or prevent development of resistance while suboptimal drug doses encourage viral replication and emergence of drug-resistance.
- Rapid availability of CMV viral load measurements as well as sensitive detection of CMV in CSF are crucial for adequate antiviral chemotherapy.
- In the presented case report, close monitoring of genotypic resistance gave an opportunity to modulate the antiviral treatment in function of the identified mutant viruses.
- In our case, the same CMV strain in different body compartments was found.



Case study II: Multidrug-resistant CMV infection in a pediatric stem

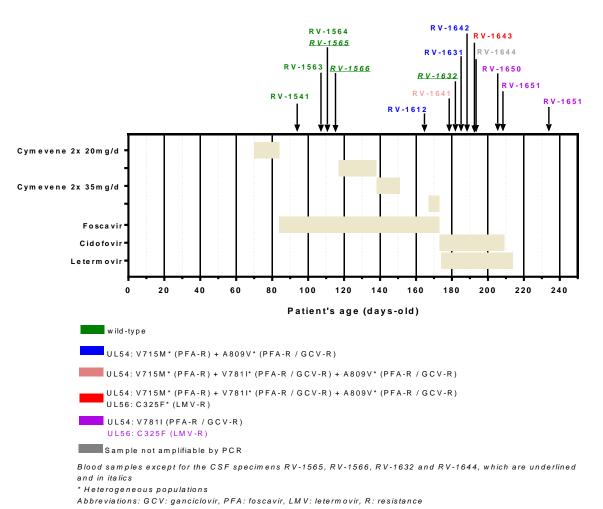
cell transplantation patient

•





Emergence of resistance to new anti-HCMV agents in an HSCT pediatric patient



| ricgavii ib | UL54 | UL56 | UL54 | UL56 |
|----------------|----------------------------|-------------|--|--------------|
| RV-1541 | Wild-type | ND | Wild-type | Wild-type |
| RV-1563 | Wild-type | ND | Wild-type | Wild-type |
| RV-1564 | Wild-type | ND | Wild-type | Wild-type |
| <u>RV-1565</u> | Wild-type | ND | ND | ND |
| <u>RV-1566</u> | Wild-type | ND | ND | ND |
| RV-1612 | V715M* A809V* | ND (| V715M: 20.5% V781I: 11.1% A809V: 23.3% | Wild-type |
| RV-1631 | V715M* A809V* | Wild-type | V715M: 20.4% V781I: 7.9% A809V: 57.6% | Wild-type |
| RV-1632 | Wild type | Wild-type | ND | ND |
| RV-1641 | V715M* V781I* A809V* | Wild-type | V715M: 12.2% V781I: 12.8% A809V: 58.1% | Wild-type |
| RV-1642 | V715M* A809V* | Wild-type (| V715M: 64.2% V781I: 6.9% A809V: 20.5% | Wild-type |
| RV-1643 | V715M* V781I* A809V* | C325F* | V715M: 19.5% V781I: 12.7% A809V: 41.8% | C325F: 26.1% |
| RV-1644 | Not amplifiable | ND | ND | ND |
| RV-1650 | V781I | C325F | V781I: 92.7% | C325F: 90.3% |
| RV-1651 | V781I | C325F | V781I: 98.5% | C325F: 98.3% |
| RV-1694 | V781I | C325F | V781I: 97.1% | C325F: 95.9% |

Sanger sequencing

RegaVir ID

NGS

- HeterogeinicityNGS advantage
- Compartmentalization

Letermovir resistance

