## **Study Synopsis**

Protocol Title	BALANCE+: A Platform Trial for Gram Negative Bloodstream Infections
<b>Protocol Number</b>	
Study Design and Phase	Perpetual multiple domain randomized controlled platform trial
Setting	International, multi-centre
Sample Size –	The initial vanguard phase will target 72 patients for most domains
BALANCE+ vanguard	parameter general process and a second process are a second process and
Sample size –	Successful domains will transition into the BALANCE + perpetual platform
BALANCE+ main platform	trial, which will have no fixed sample size
Platform Entry Criteria	Inclusion criteria:
	admitted to a participating hospital
	positive blood culture with Gram negative (GN) bacterium
	Exclusion criteria:
	patient's goals of care are for palliation with no active treatment
	<ul> <li>moribund patient, not expected to survive &gt; 72 hours</li> </ul>
	monitorial patients, not expected to survive 7.2 nours
	There are additional domain-specific inclusion/exclusion criteria
Domains	De-escalation versus no de-escalation domain
<ul><li>intervention</li></ul>	No de-escalation
arms	o continue on the same empiric GN antibiotic(s) being
	used prior to blood culture finalization
	<ul> <li>companion antibiotics in a combination regimen (eg,</li> </ul>
	vancomycin, azithromycin, aminoglycoside or
	metronidazole) can be discontinued at discretion of
	clinical team
	<ul> <li>if patient has a syndrome requiring prolonged duration</li> </ul>
	(> 14 days) de-escalation is allowable after day 14
	De-escalation
	<ul> <li>empiric GN antibiotic(s) switched to narrower spectrum agent to which the blood culture isolate is susceptible</li> </ul>
	(within 24 hours)
	o companion antibiotics in a combination regimen can be
	discontinued at discretion of clinical team (and will be
	encouraged to do so in this arm if not needed for another indication)
	Oral beta-lactam versus non beta-lactam domain
	<ul> <li>Non-beta-lactam arm: an oral fluoroquinolone (ciprofloxacin,</li> </ul>
	moxifloxacin or levofloxacin) or trimethoprim-sulfamethoxazole

	<ul> <li>Beta-lactam arm: an oral beta-lactam agent including, but not limited to, amoxicillin, amoxicillin-clavulanate, cephalexin, cefadroxil, or cefixime</li> </ul>
	<ul> <li>Central vascular catheter retention versus replacement domain</li> <li>Central vascular catheter retention arm: original line retained until non-functional, or no longer needed</li> <li>Central vascular catheter replacement arm: replace the vascular catheter as soon as possible and within a maximum of 72 hours from blood culture finalization</li> </ul>
	<ul> <li>Cephalosporin arm: participants will be treated with ceftriaxone (at standard doses) during intravenous treatment, with oral step-down allowed to any susceptible agent</li> <li>Carbapenem arm: participants will be treated with a carbapenem (at standard doses) during intravenous treatment, with oral step-down allowed to any susceptible agent</li> </ul>
Primary Feasibility	<ul> <li>Follow-up blood culture domain</li> <li>Follow-up blood culture arm: participants will undergo routine repeat blood culture collection (at least one blood culture set)         4±1d from the calendar date of the index positive blood culture collection     </li> <li>No follow-up blood culture arm: participants will undergo no routine repeat blood culture collection between 4±1d from the calendar date of the index positive blood culture collection</li> <li>Domain-specific recruitment rate</li> </ul>
Outcomes of BALANCE+ vanguard	Domain-specific protocol adherence
Primary Outcome of BALANCE+ domains	<ul> <li>De-escalation versus no de-escalation domain</li> <li>Patient-centered, ordinal Desirability of Outcome Ranking (DOOR) outcome: (dead at 90 days) &lt; (alive at 90 days with reinfection and readmission) &lt; (alive at 90 days with neither reinfection nor readmission)</li> <li>tie-breaker within ordinal levels: new antimicrobial resistance (AMR) colonization or infection from routine cultures</li> <li>Oral beta-lactam versus non beta-lactam domain</li> <li>ordinal DOOR outcome: (dead at 90 days) &lt; (alive at 90 days with reinfection and readmission) &lt; (alive at 90 days with neither reinfection nor readmission)</li> </ul>



	<ul> <li>tie-breaker within ordinal levels: new AMR colonization or infection from routine cultures</li> <li>Central vascular catheter retention versus replacement domain</li> <li>ordinal DOOR outcome: (dead at 90 days) &lt; (alive at 90 days with reinfection and readmission) &lt; (alive at 90 days with neither reinfection nor readmission)</li> <li>no tie-breaker</li> <li>Low-risk AmpC domain</li> <li>ordinal DOOR outcome: (dead at 90 days) &lt; (alive at 90 days</li> </ul>
	with reinfection and readmission) < (alive at 90 days with reinfection or readmission) < (alive at 90 days with neither reinfection nor readmission)  • tie-breaker within ordinal levels: new AMR colonization or infection from routine cultures  Follow-up blood culture domain  • ordinal DOOR outcome: (dead at 90 days) < (alive at 90 days with reinfection and readmission) < (alive at 90 days with neither
	reinfection nor readmission)  • no tie-breaker
Secondary Outcomes of BALANCE+	<ul> <li>90-day mortality</li> <li>90-day reinfection</li> <li>90-day all cause readmission</li> <li>90-day AMR colonization/infection</li> <li>90-day Clostridioides difficile infection (CDI)</li> <li>30-day mortality</li> <li>60-day mortality</li> <li>additional domain-specific secondary outcomes</li> </ul>
Statistical Analysis BALANCE+ vanguard	<ul> <li>descriptive point estimates and 95% confidence intervals for domain-specific recruitment rates and protocol adherence</li> <li>analyzed everall and by participating site</li> </ul>
Statistical Analysis BALANCE+ Main Platform	<ul> <li>analyzed overall and by participating site</li> <li>regular Bayesian interim analyses with uninformative priors</li> <li>conducted at every 500th BALANCE+ platform enrolment</li> <li>domains closed only if they meet pre-specified, stringent decision criteria for stopping based on superiority, non-inferiority or futility</li> </ul>