

Patient Information					
First Name	MI	Last Name	Medical Record #	DOB	Sex <input type="radio"/> Male <input type="radio"/> Female
Address		City	State	Postal Code	Country
					Primary Phone

Patient Medical History	
Primary ICD-10	Stage
Prior / Current Therapies (Optional)	

Diagnosis		
<input type="radio"/> Colorectal Carcinoma	<input type="radio"/> NSCLC	
<input type="radio"/> Breast	<input type="radio"/> Prostate	<input type="radio"/> Melanoma
<input type="radio"/> Ovarian	<input type="radio"/> Other _____	

Disease Status (Select all that apply)	
<input type="radio"/> Metastatic	<input type="radio"/> Recurrent
<input type="radio"/> Refractory	<input type="radio"/> Relapse
<input type="radio"/> None	<input type="radio"/> Progression

Attachments	
<input type="checkbox"/> Copy of recent pathology/cytology reports including (if available), CBC/differential, BMA differential, FAB classification.	<input type="checkbox"/> Test results from all other Molecular Diagnostic Assays by FISH, IHC, or other genetic assays, e.g., ER, PR, HER2, EGFR, KRAS, etc.

Ordering Physician Information					
Facility Name			Physician Name		
Address			Phone	Fax	
City	State	Postal Code	Email		
Is the facility a hospital, hospital outpatient department, critical access hospital or ambulatory surgical center? (see back)			<input type="radio"/> No	<input type="radio"/> Yes →	If yes, what is the facility's network status with the patient's insurance plan?
			<input type="radio"/> In-Network	<input type="radio"/> Out-of-Network	<input type="radio"/> Unknown

Genomic Test	Description	Accepted Specimen Type
<input type="checkbox"/> Aventa FusionPlus Test	CLIA-certified, test with a clinical report covering 361 genes implicated in solid tumors across multiple tumor types.	<input type="checkbox"/> FFPE Tissue Block <input type="checkbox"/> 10 x 5 µm FFPE Tissue Sections

Specimen Retrieval											
Submitting Pathologist Name	Pathology Lab Name	Email	Phone	Fax							
<input type="checkbox"/> I am requesting a specific specimen <table border="1"> <tr> <td>Collection Date (MM/DD/YYYY)</td> <td>Specimen ID</td> <td>Site of Biopsy</td> </tr> <tr> <td></td> <td></td> <td></td> </tr> </table>					Collection Date (MM/DD/YYYY)	Specimen ID	Site of Biopsy				<b>Shipment:</b> <input type="checkbox"/> I will arrange for specimen shipment <input type="checkbox"/> Contact the pathology lab to obtain specimen
Collection Date (MM/DD/YYYY)	Specimen ID	Site of Biopsy									
<input type="checkbox"/> I will let the pathologist choose the specimen <input type="checkbox"/> I am providing FFPE block return address on back of form											

Insurance Billing Information					
<input type="checkbox"/> Medicare – Part B:	<input type="checkbox"/> *ABN Attached (If required, see back)	Medicare Policy ID	*Patient status at the time of specimen collection:	<input type="checkbox"/> Office (non-hospital) <input type="checkbox"/> Outpatient <input type="checkbox"/> Inpatient: →	<input type="checkbox"/> Not yet discharged Discharge date
<input type="checkbox"/> Insurance:	Plan Name	Policy #	Group #	Prior Authorization #	
<input type="checkbox"/> Self-Pay:	Contact Name	Email	Phone		
<input type="checkbox"/> Hospital / Institution:	<input type="checkbox"/> Same as treating physician	Address			
	City	State	Postal Code	Fax	

Submission Checklist	Physician Signature and Letter of Medical Necessity				
<input type="checkbox"/> Demographic / Face Sheet <input type="checkbox"/> Most recent office note <input type="checkbox"/> Pathology Report <input type="checkbox"/> Copy of insurance cards	My signature certifies that I have determined that the test(s) being ordered is medically necessary for the patient, certifies that the results of this test will inform the patient's ongoing treatment plan, and certifies that I am the patient's treating physician. I have explained to the patient the nature and purpose of the test(s) to be performed and have obtained informed consent, to the extent required under applicable law, to permit Aventa Genomics, or any laboratory with which Aventa Genomics is contracted, to (a) perform the test(s) specified herein, (b) analyze and report on other genetic information generated during the testing process or conduct additional analyses of the patient's sample for future diagnostic or monitoring use, (c) retain the test results and tissues, cells, and genetic material, including DNA and RNA information generated during the testing process, for an indefinite period for internal quality assurance/operations purposes, (d) remove information that directly identifies the patient from the test results, tissues, cells, and genetic material, including DNA and RNA information generated during the testing process, and use or disclose such information and materials for future unspecified research or other purposes, and (e) release the test results and related patient information to the patient's third-party payer as needed for reimbursement purposes.				
	<table border="1"> <tr> <td>Ordering Physician Signature</td> <td>Date</td> </tr> <tr> <td></td> <td></td> </tr> </table>	Ordering Physician Signature	Date		
Ordering Physician Signature	Date				

FFPE Block Return Information					
Return address		City	State	Postal Code	Country
Email		Phone		Fax	

Additional Case Information

Test Description	Sample Requirements
<p>The Aventa FusionPlus test utilizes a method known as HiC sequencing which is designed specifically to capture a genome's sequence and structure (three-dimensional conformation). FFPE tissue sections are dewaxed and rehydrated. Then the cross-linked chromatin is digested using a restriction enzyme (RE) cocktail. The 5'-overhangs are then filled in with a biotinylated nucleotide. Next, spatially proximal digested ends of DNA are ligated, capturing the sequence and structure of the genome. The ligated DNA is then purified, producing pure proximally-ligated DNA. The proximally-ligated DNA is then fragmented, and the biotinylated fragments are enriched. DNA libraries are then prepared from these enriched libraries. Finally, libraries are sequenced in a "paired-end" mode.</p> <p><b>Secondary Analysis Methods:</b>            The resulting data is processed using the Arima-SV Pipeline. The pipeline is used for calling and visualizing Structural Variants (SV). This pipeline preprocesses the data using HiCUP (Wingett et al. 2015) and calls SV's using hic_breakfinder (Dixon et al. 2018). The SV's are manually curated and processed to create a single VCF file that is directly ingested into CGW.</p>	<p>This testing service requires 10 x 5µm FFPE tissue sections, scrolls or tissue block and an H&amp;E stained tissue section.</p> <p><b>For information on ICD codes</b>            Visit this website: <a href="https://icd10cmtool.cdc.gov/">https://icd10cmtool.cdc.gov/</a></p>